

Network Systems  
Science & Advanced  
Computing  
Biocomplexity Institute  
& Initiative  
University of Virginia

# Estimation of COVID-19 Impact in Virginia

January 13<sup>th</sup>, 2021

(data current to January 11<sup>th</sup>-12<sup>th</sup>)

Biocomplexity Institute Technical report: TR 2021-004



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**BIOCOMPLEXITY** INSTITUTE

[biocomplexity.virginia.edu](https://biocomplexity.virginia.edu)

# About Us

- Biocomplexity Institute at the University of Virginia
  - Using big data and simulations to understand massively interactive systems and solve societal problems
- Over 20 years of crafting and analyzing infectious disease models
  - Pandemic response for Influenza, Ebola, Zika, and others



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# Overview

- **Goal:** Understand impact of COVID-19 mitigations in Virginia
- **Approach:**
  - Calibrate explanatory mechanistic model to observed cases
  - Project based on scenarios for next 4 months
  - Consider a range of possible mitigation effects in "what-if" scenarios
- **Outcomes:**
  - Ill, Confirmed, Hospitalized, ICU, Ventilated, Death
  - Geographic spread over time, case counts, healthcare burdens

# Key Takeaways

Projecting future cases precisely is impossible and unnecessary.  
Even without perfect projections, we can confidently draw conclusions:

- **Case rate growth in Virginia continues to surge along with the nation, but remains below national average**
- VA mean weekly incidence (60/100K) up (from 52) as national levels also rebound (to 67/100K from 60/100K); Virginia records highest daily case rate in past week
- Projections are mostly up across commonwealth
- Recent updates:
  - Modified scenarios to be based on past control levels (best and fatigued)
  - Planning scenarios start changing transmission rates at end of the month (Jan 30)
  - Refined vaccination schedule to account for partial protection from first dose
  - Adjusted “rescaling” method and data sources to better accommodate recent trends
- The situation is changing rapidly. Models will be updated regularly.

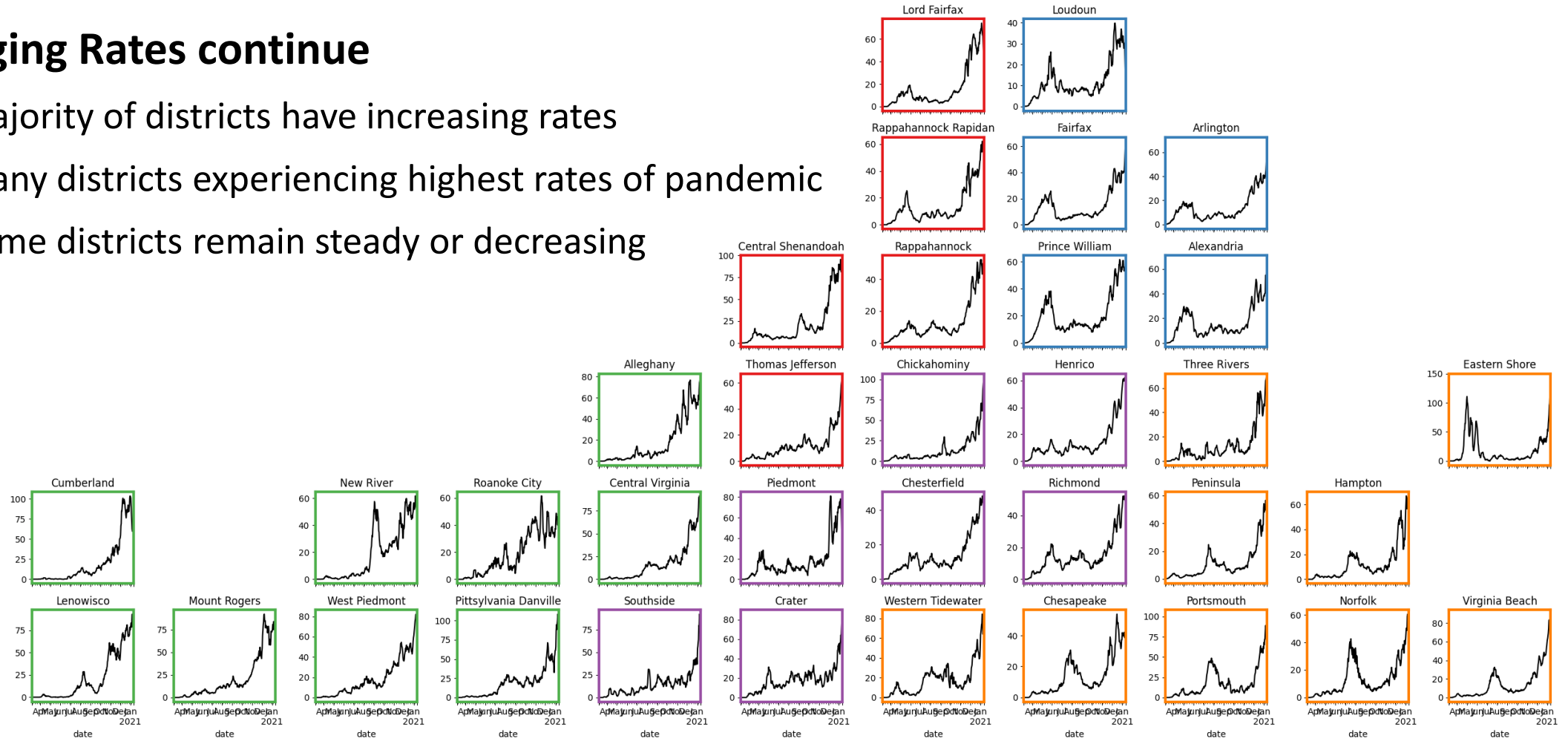
# Situation Assessment

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# Case Rate (per 100k) by VDH District

## Surging Rates continue

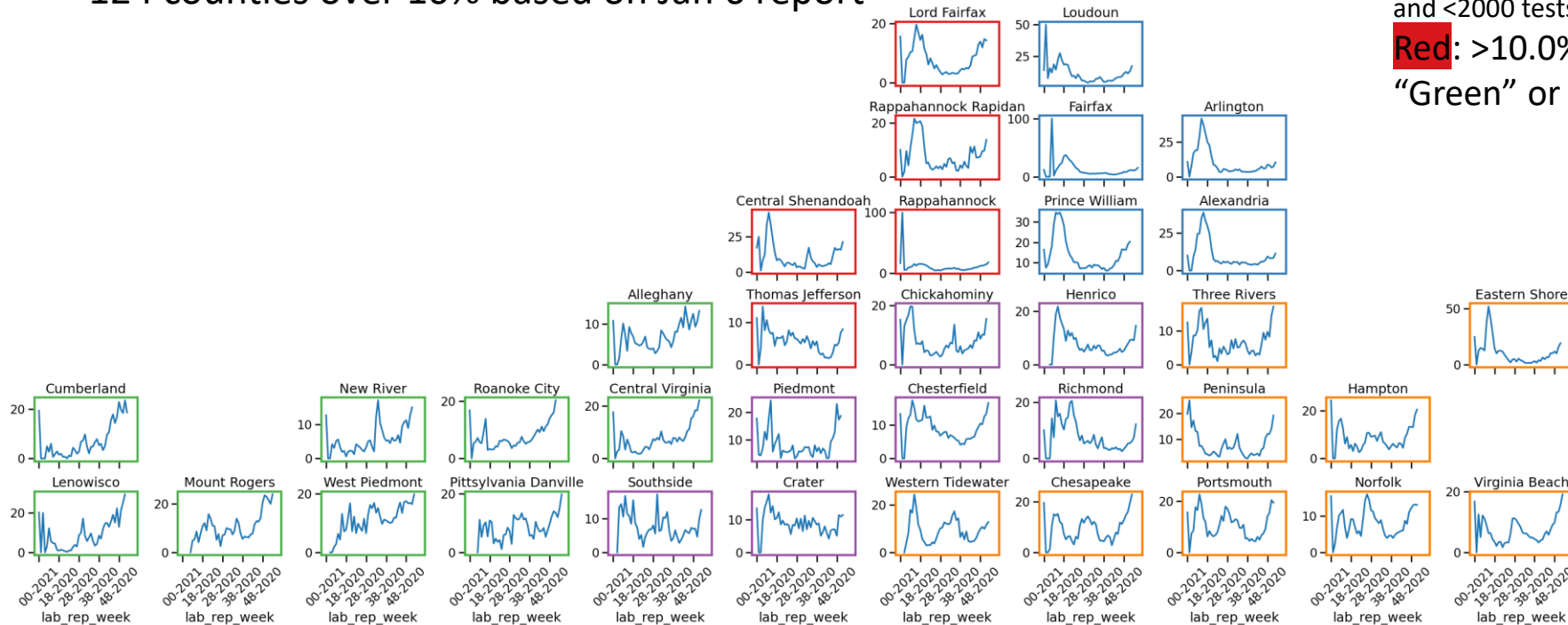
- Majority of districts have increasing rates
- Many districts experiencing highest rates of pandemic
- Some districts remain steady or decreasing



# Test Positivity by VDH District

## Weekly changes in test positivity by district

- Increasing levels in many districts throughout the commonwealth with many districts above 10% for several weeks
- 124 counties over 10% based on Jan 6 report

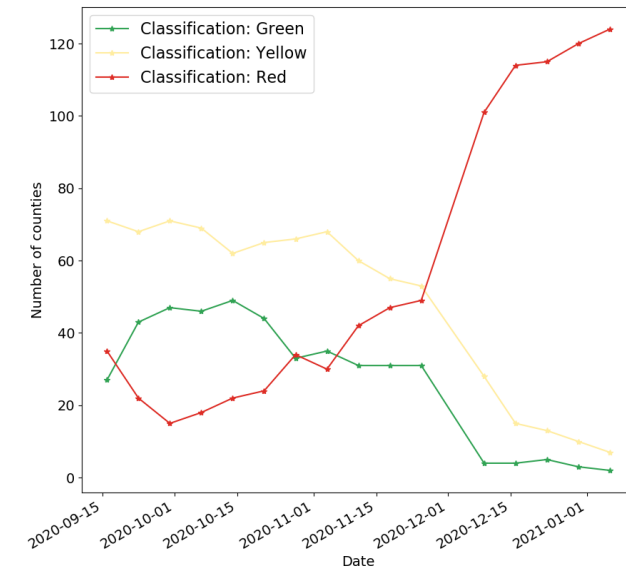


## County level test positivity rates for RT-PCR tests.

**Green:** Test positivity <5.0%  
(or with <20 tests in past 14 days)

**Yellow:** Test positivity 5.0%-10.0% (or with <500 tests and <2000 tests/100k and >10% positivity over 14 days)

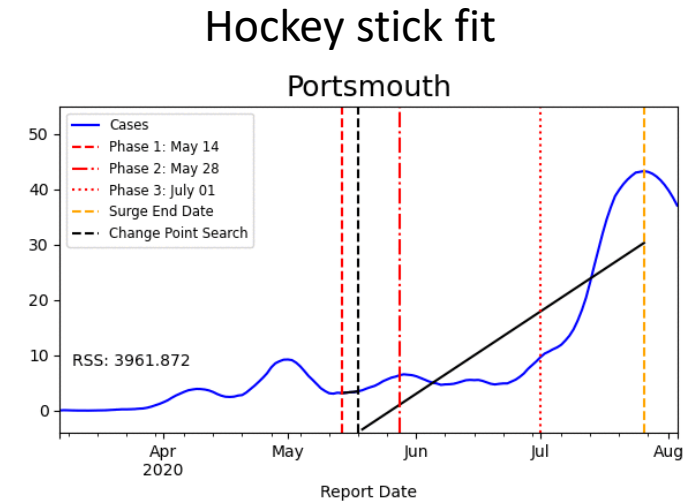
**Red:** >10.0% and not meeting the criteria for “Green” or “Yellow”



# District Trajectories

**Goal:** Define epochs of a Health District's COVID-19 incidence to characterize the current trajectory

**Method:** Find recent peak and use hockey stick fit to find inflection point afterwards, then use this period's slope to define the trajectory



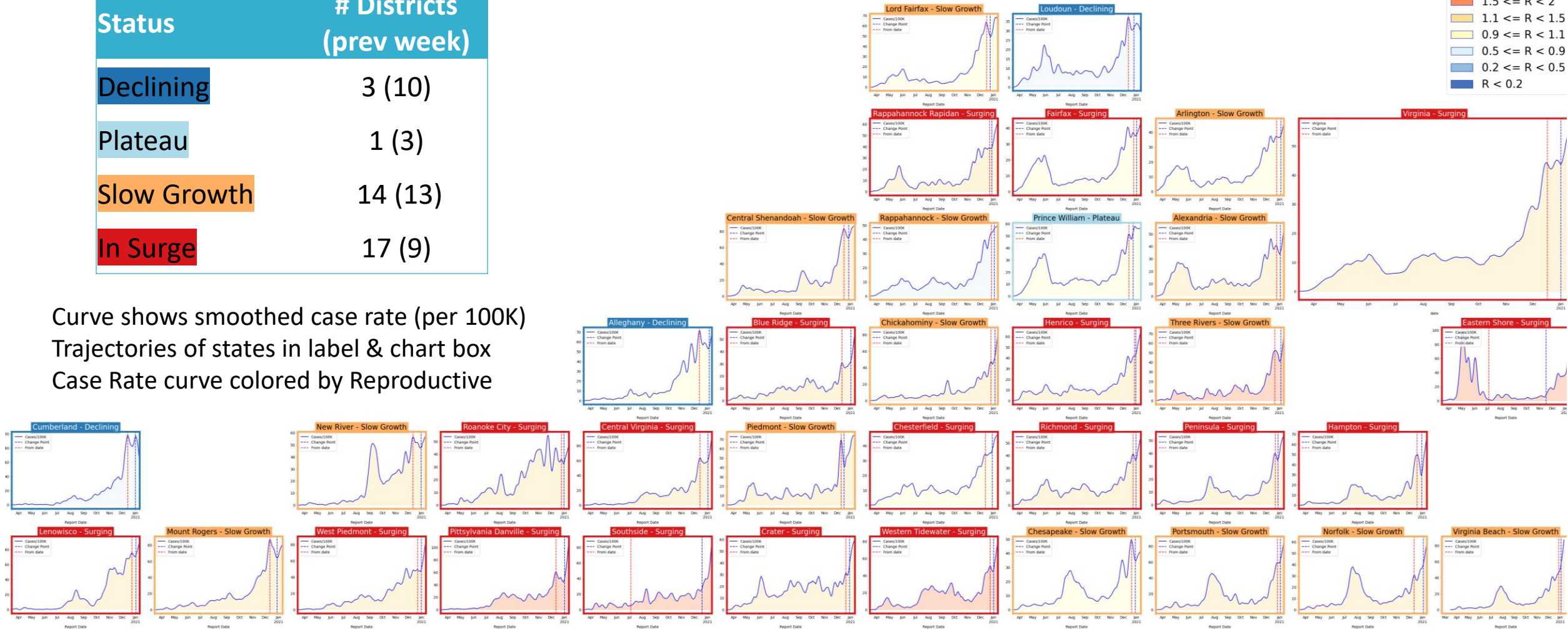
Trajectory	Description	Weekly Case Rate (per 100K) bounds	# Districts (prev week)
<b>Declining</b>	Sustained decreases following a recent peak	below -0.9	3 (10)
<b>Plateau</b>	Steady level with minimal trend up or down	above -0.9 and below 0.5	1 (3)
<b>Slow Growth</b>	Sustained growth not rapid enough to be considered a Surge	above 0.5 and below 2.5	14 (13)
<b>In Surge</b>	Currently experiencing sustained rapid and significant growth	2.5 or greater	17 (9)



# District Trajectories – last 10 weeks

Status	# Districts (prev week)
Declining	3 (10)
Plateau	1 (3)
Slow Growth	14 (13)
In Surge	17 (9)

Curve shows smoothed case rate (per 100K)  
Trajectories of states in label & chart box  
Case Rate curve colored by Reproductive



# Estimating Daily Reproductive Number

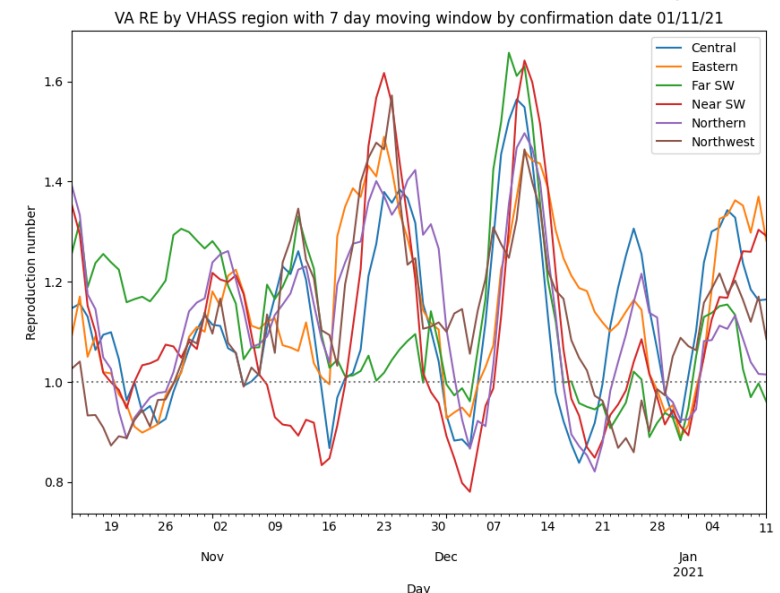
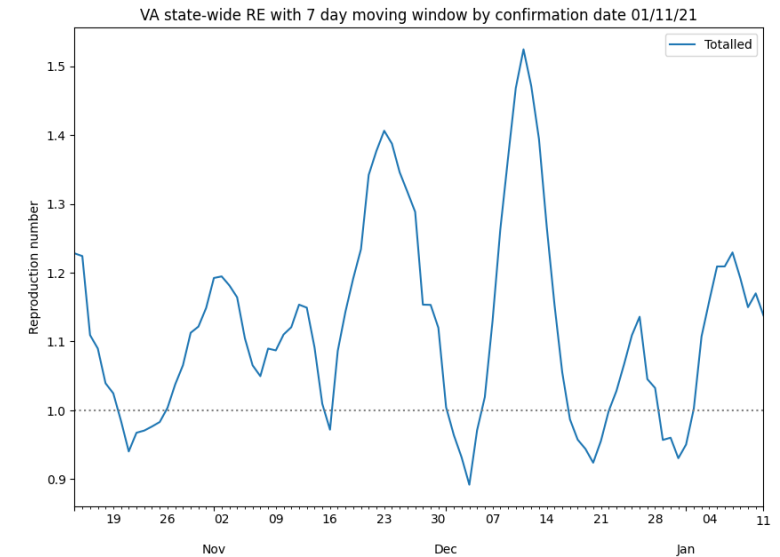
## Jan 11<sup>th</sup> Estimates

Region	Date Confirmed $R_e$	Date Confirmed Diff Last Week
State-wide	1.138	-0.021
Central	1.164	-0.136
Eastern	1.282	0.096
Far SW	0.960	-0.177
Near SW	1.291	0.167
Northern	1.015	-0.069
Northwest	1.086	-0.100

### Methodology

- Wallinga-Teunis method (EpiEstim<sup>1</sup>) for cases by **confirmation date**
- Serial interval: 6 days (2 day std dev)
- Using Confirmation date since due to increasingly unstable estimates from onset date due to backfill

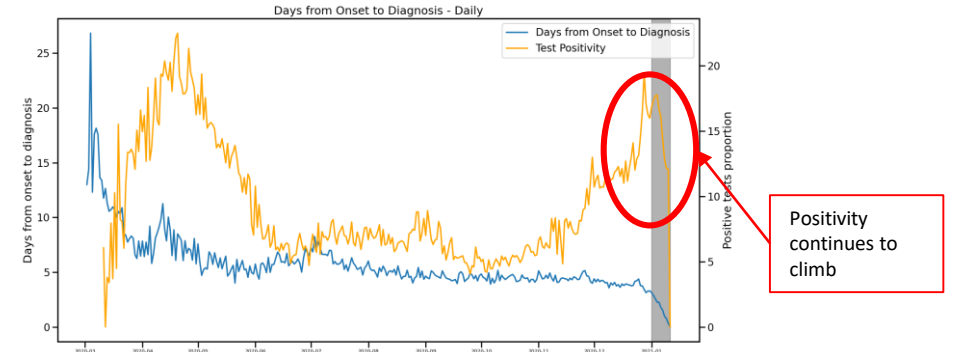
1. Anne Cori, Neil M. Ferguson, Christophe Fraser, Simon Cauchemez. A New Framework and Software to Estimate Time-Varying Reproduction Numbers During Epidemics. American Journal of Epidemiology, Volume 178, Issue 9, 1 November 2013, Pages 1505–1512, <https://doi.org/10.1093/aje/kwt133>



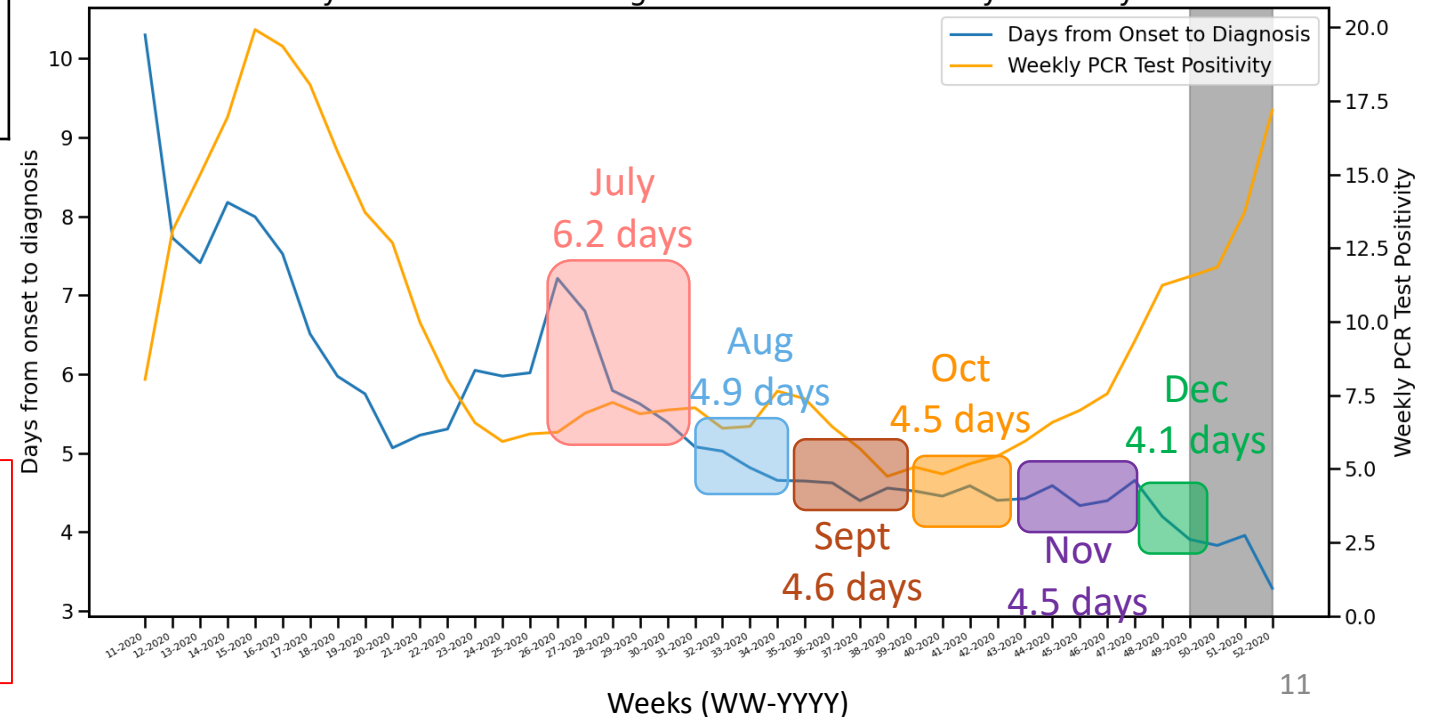
# Changes in Case Detection

Timeframe (weeks)	Mean days	% difference from overall mean
April (13-16)	7.8	44%
May (17-21)	5.7	6%
June (22-25)	5.8	8%
July (26-30)	6.2	14%
Aug (31-34)	4.9	-9%
Sept (35-38)	4.6	-16%
Oct (39-43)	4.5	-17%
Nov (44-47)	4.5	-17%
Dec (48-49)	4.1	-25%
Overall (13-49)	5.4	--

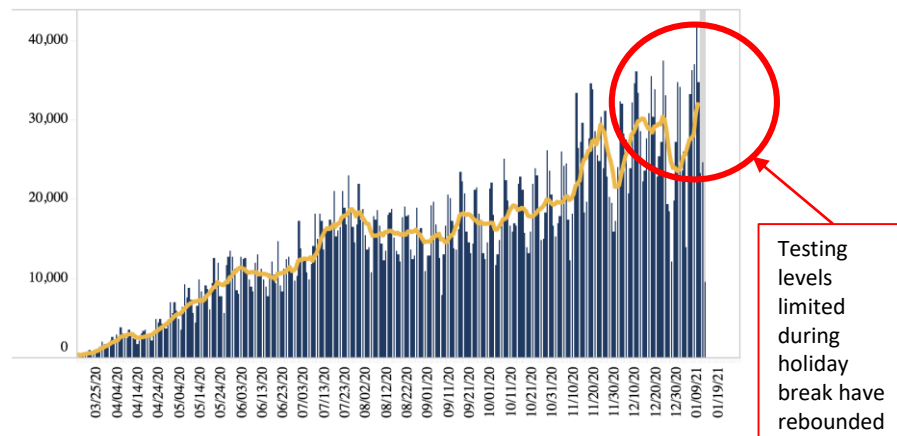
Test positivity vs. Onset to Diagnosis



Days from Onset to Diagnosis and Test Positivity - Weekly

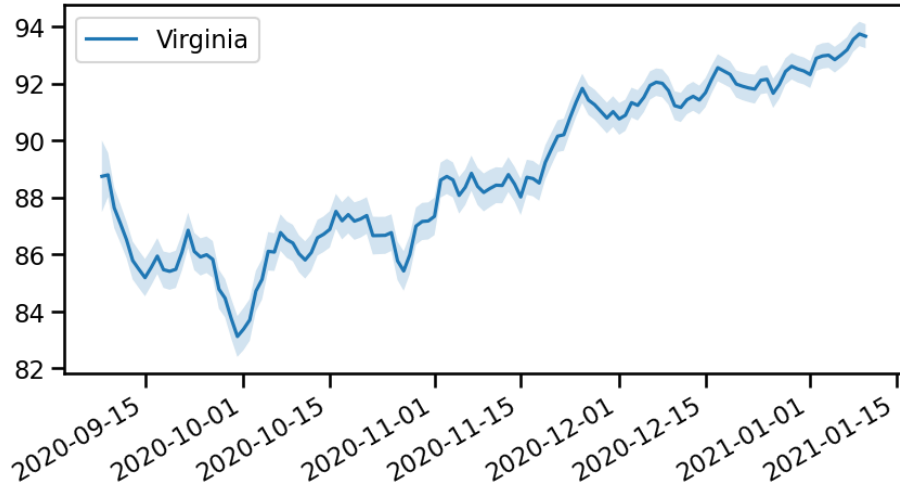


Number of Testing Encounters by Lab Report Date - All Health Districts, PCR Only



Accessed 8:30am January 13, 2021  
<https://www.vdh.virginia.gov/coronavirus/>

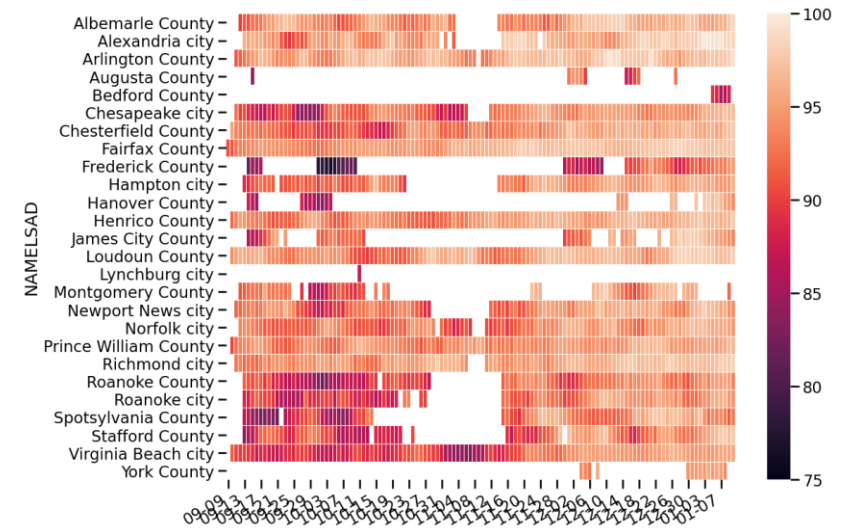
# Mask usage in Virginia



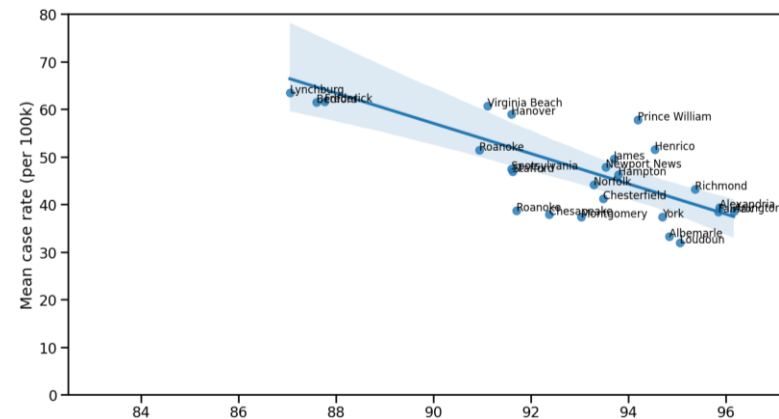
State level mask usage as reported via Facebook surveys has shown steady increase over past three months

- ~88% (early Nov) to ~94% (mid Jan)
- Some variance across the commonwealth
- ~3000 daily responses from VA

Data Source: <https://covidcast.cmu.edu>



Some county level fluctuations since beginning of Sept., though data quality may be affected by sample sizes.

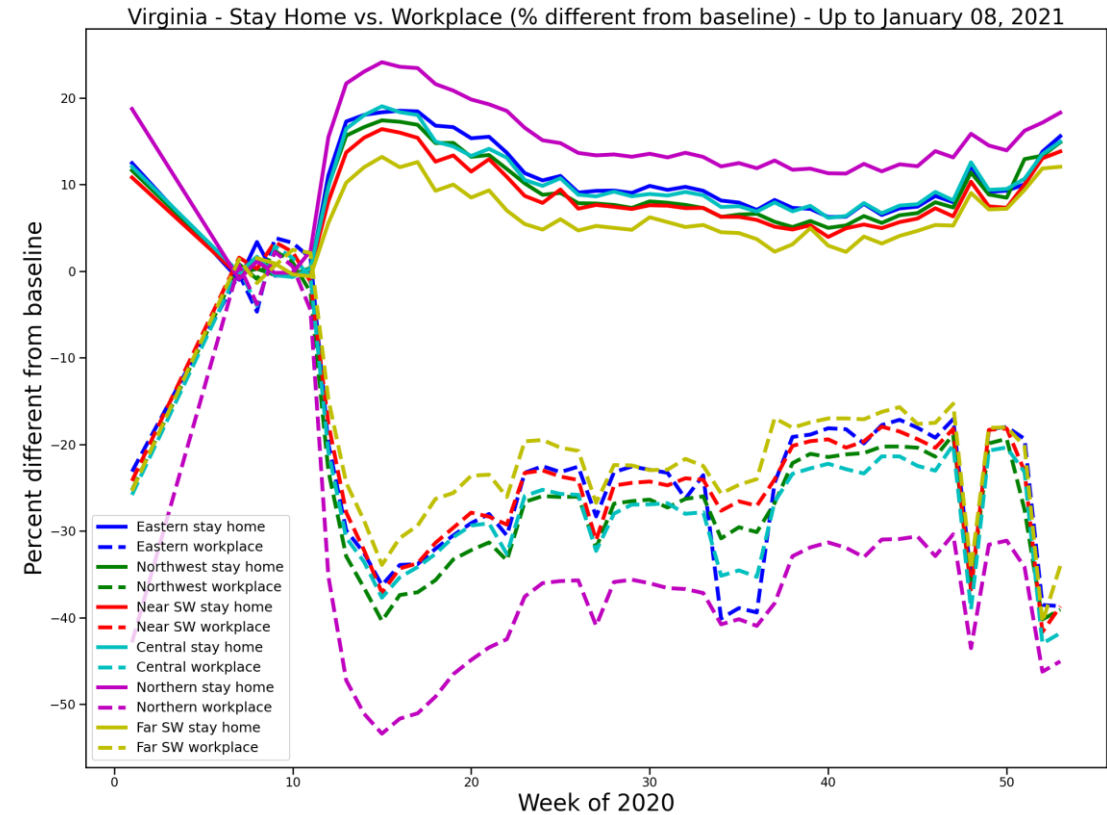
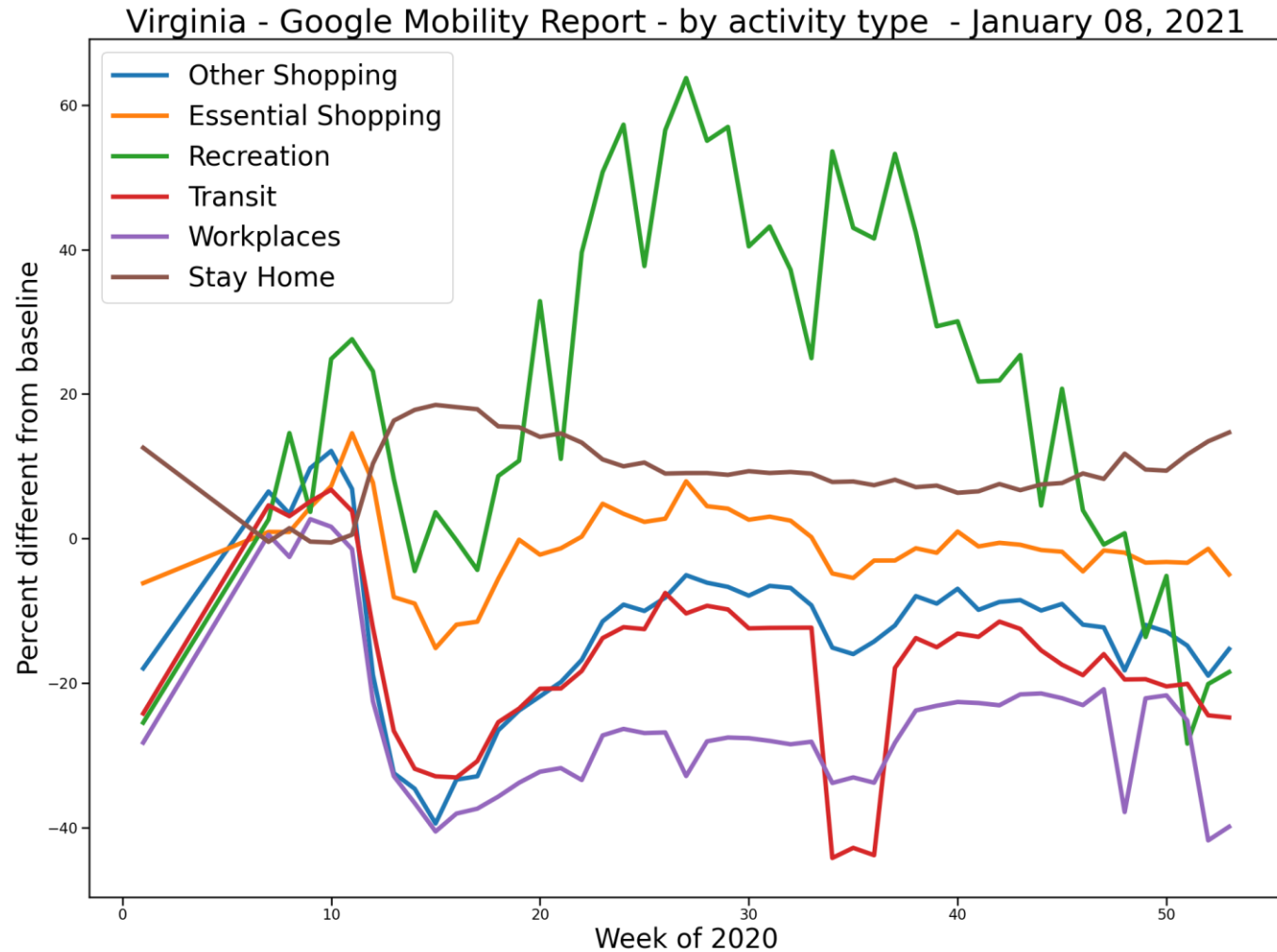


Correlations seen among VA counties between mask use and case rate are now stronger due to surging growth

Slope: - 3.2; for every % we see a ~3/100K case rate difference



# Mobility Shifts



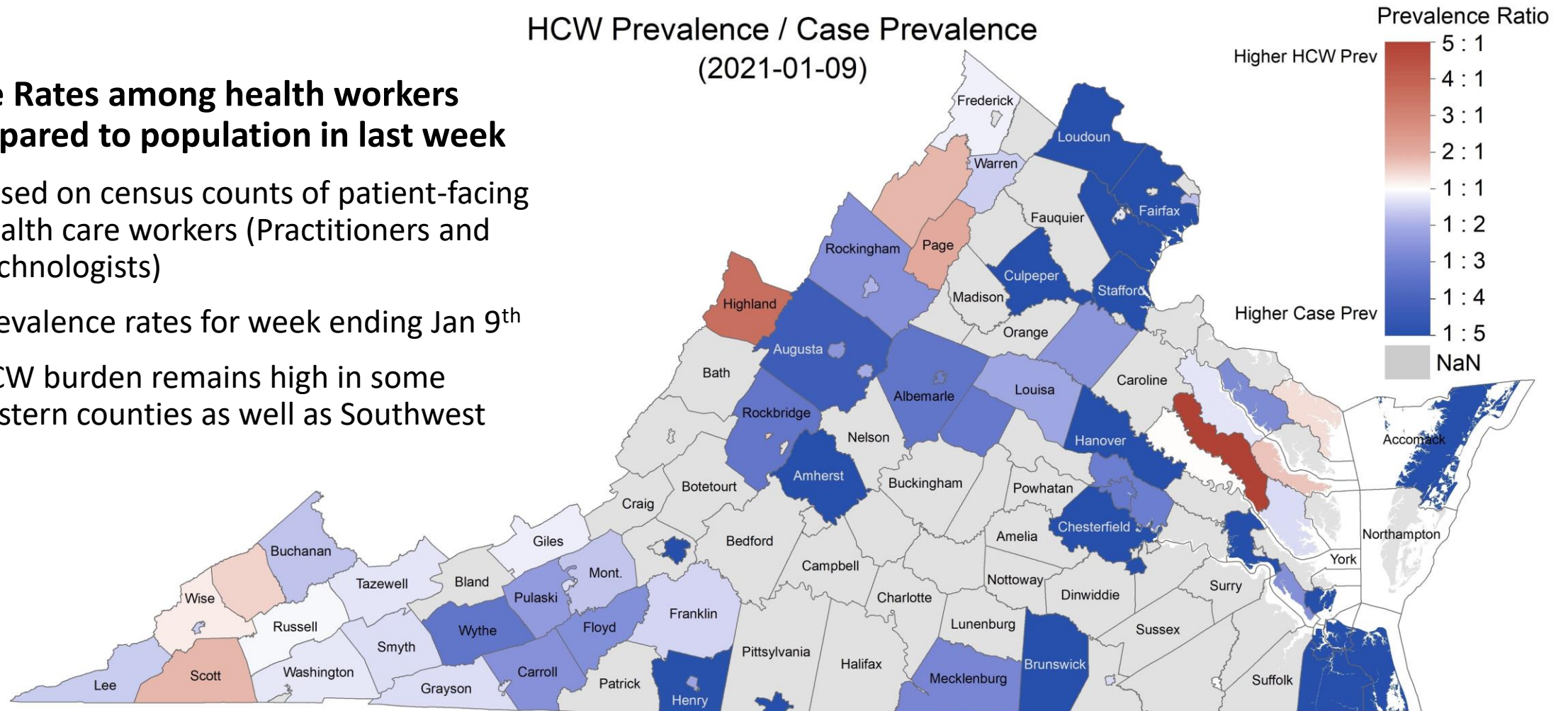
**Staying Home is on the rise, and recreation down**

- Uniform across regions
- Home activities don't preclude transmission in small gatherings

# Health Care Worker Prevalence (per 100K)

## Case Rates among health workers compared to population in last week

- Based on census counts of patient-facing health care workers (Practitioners and Technologists)
- Prevalence rates for week ending Jan 9<sup>th</sup>
- HCW burden remains high in some Eastern counties as well as Southwest

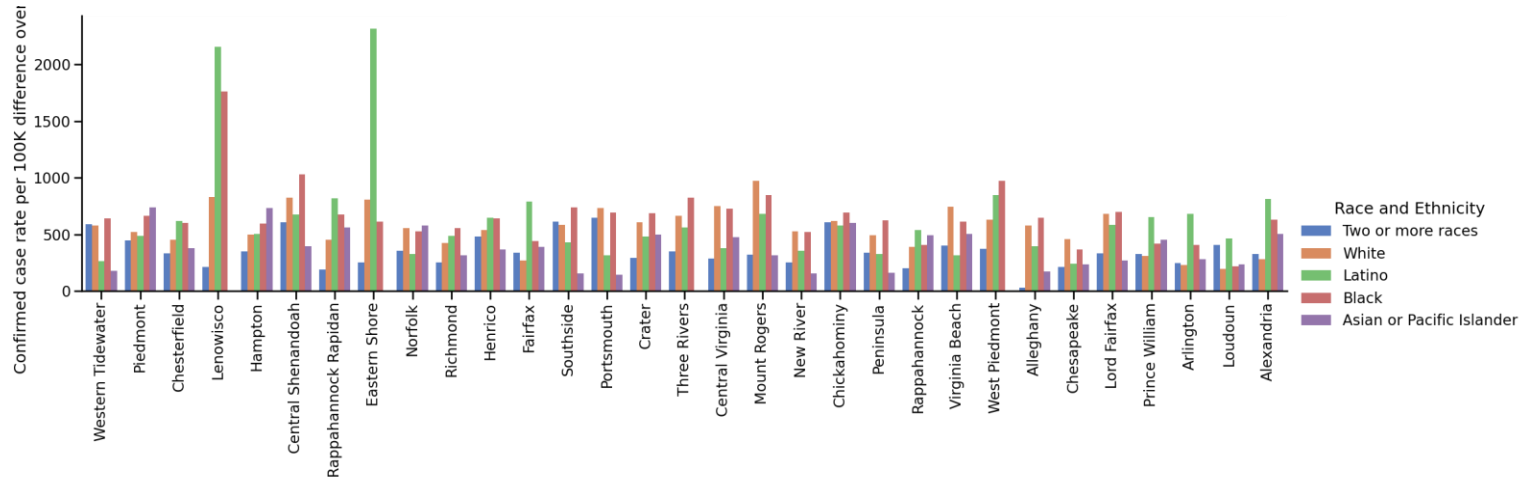


# Race and Ethnicity – Recent Rate Changes (per 100K)

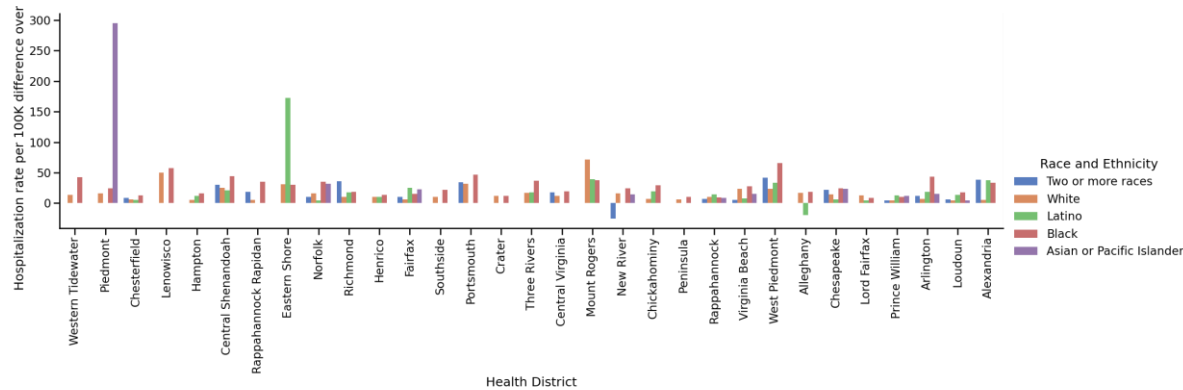
## Recent Changes in Race and Ethnicity Rates (per 100k)

- Two week change in population level rates
- Black, Latinx and 2 or more races populations have much higher changes in rates; disparity is more pronounced in some districts than others
- Based on 2019 census race-ethnicity data by county

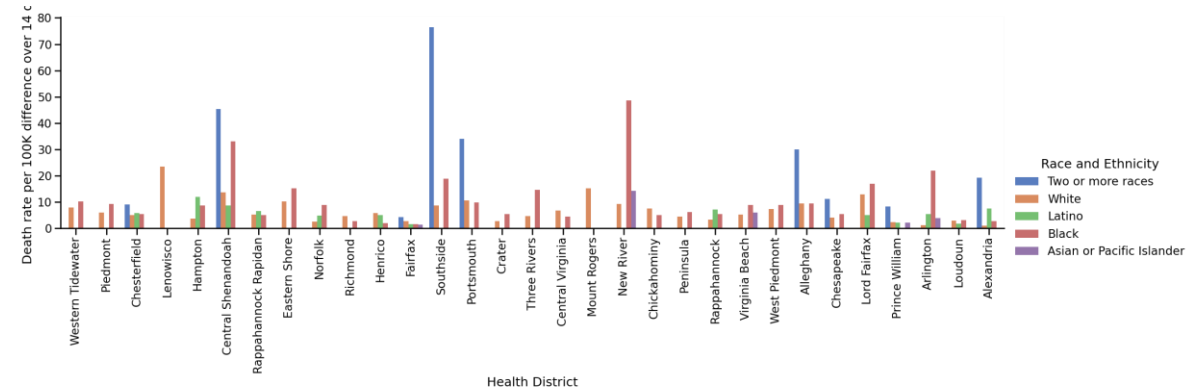
## Case Rate



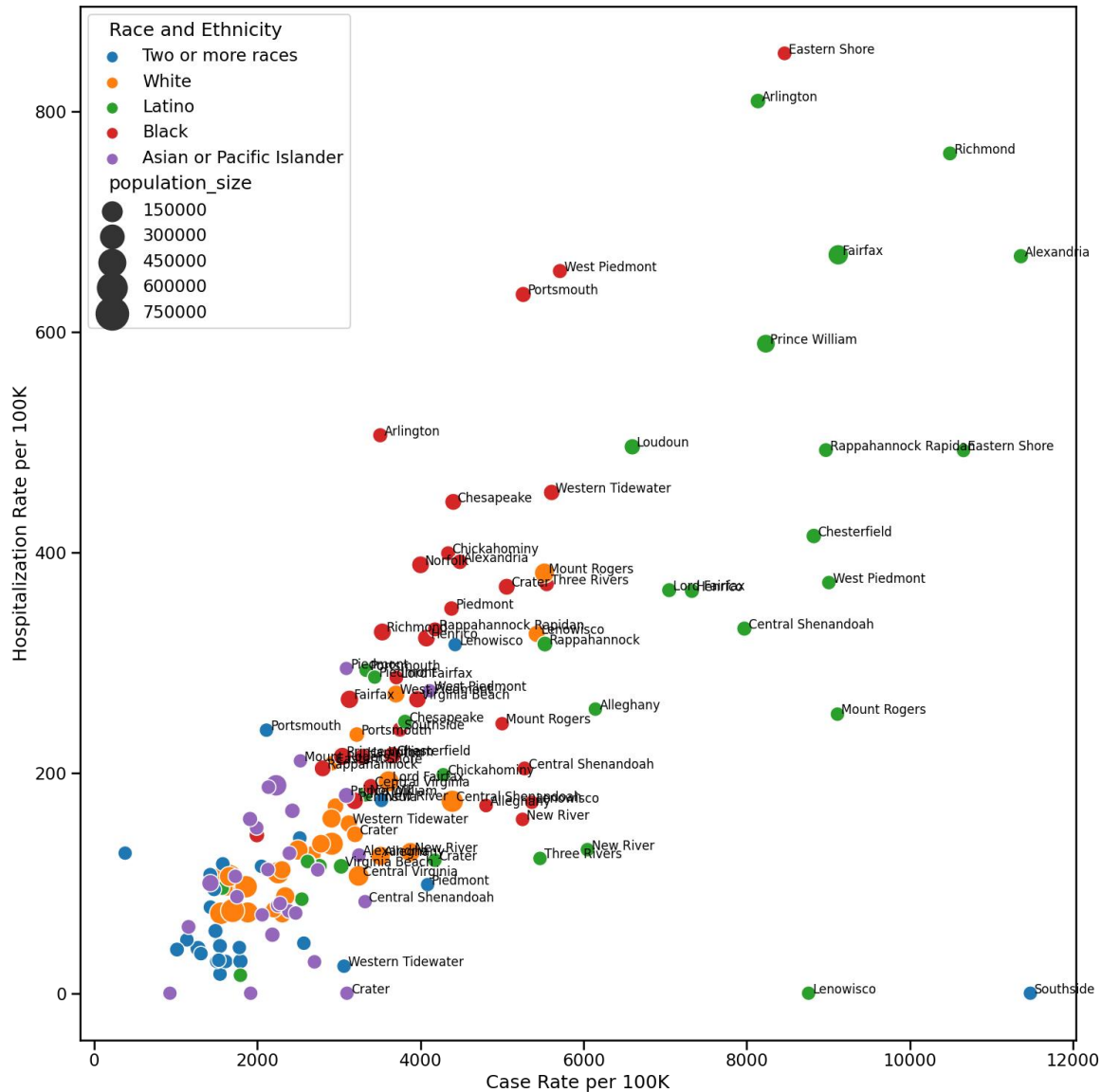
## Hospitalization Rate



## Death Rate



# Race and Ethnicity cases per 100K



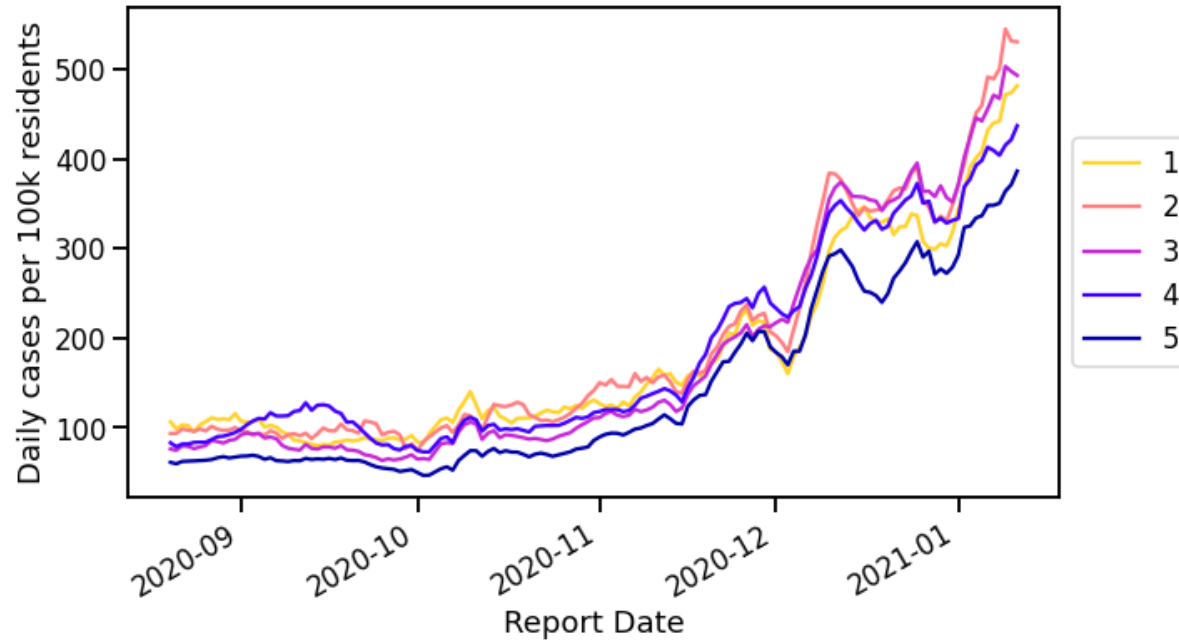
## Rates per 100K of each Racial-Ethnic population by Health District

- Each Health District's Racial-Ethnic population is plotted by their Hospitalization and Case Rate
- Points are sized based on their overall population size
- Overlapping labels removed for clarity



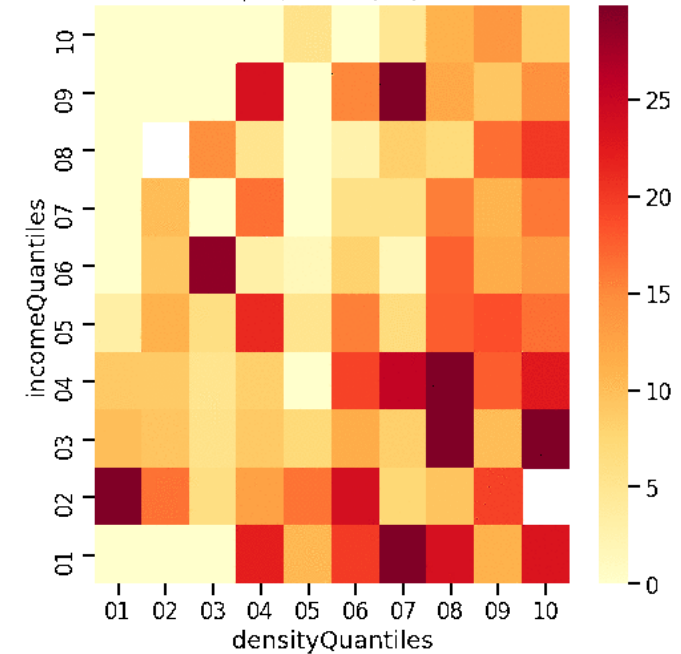
# Impact across Density and Income

VDH 7-day moving average rate of new COVID-19 cases by zip code  
average household income (dollars/ household years) quantile



All zip codes show back into growth, wealthiest zip code now lags the rest significantly

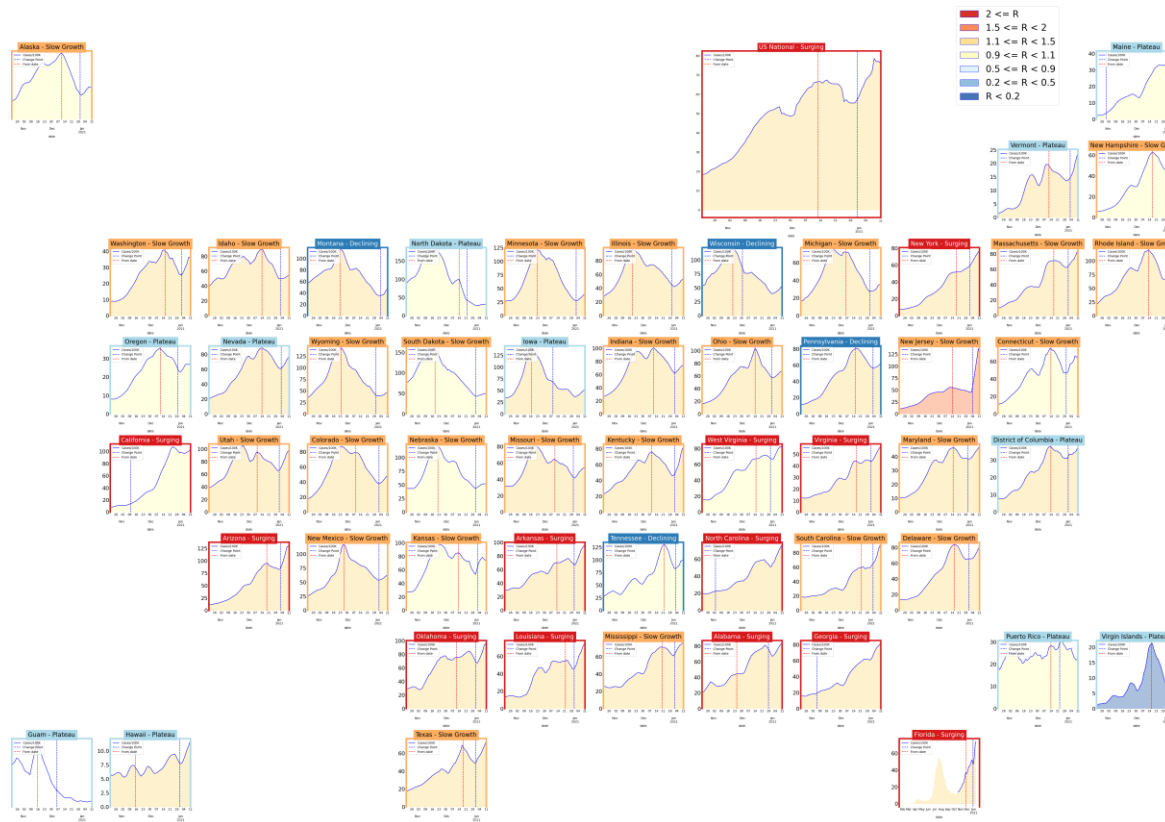
VDH mean cases per 100k by zip code population density (person/ sq mile)  
and average household income (dollars/ household years) quantiles  
08/08/20 - 08/14/20



Full evolution of pandemic, shows shifts from denser and wealthier zip codes to poorer and less dense zip codes

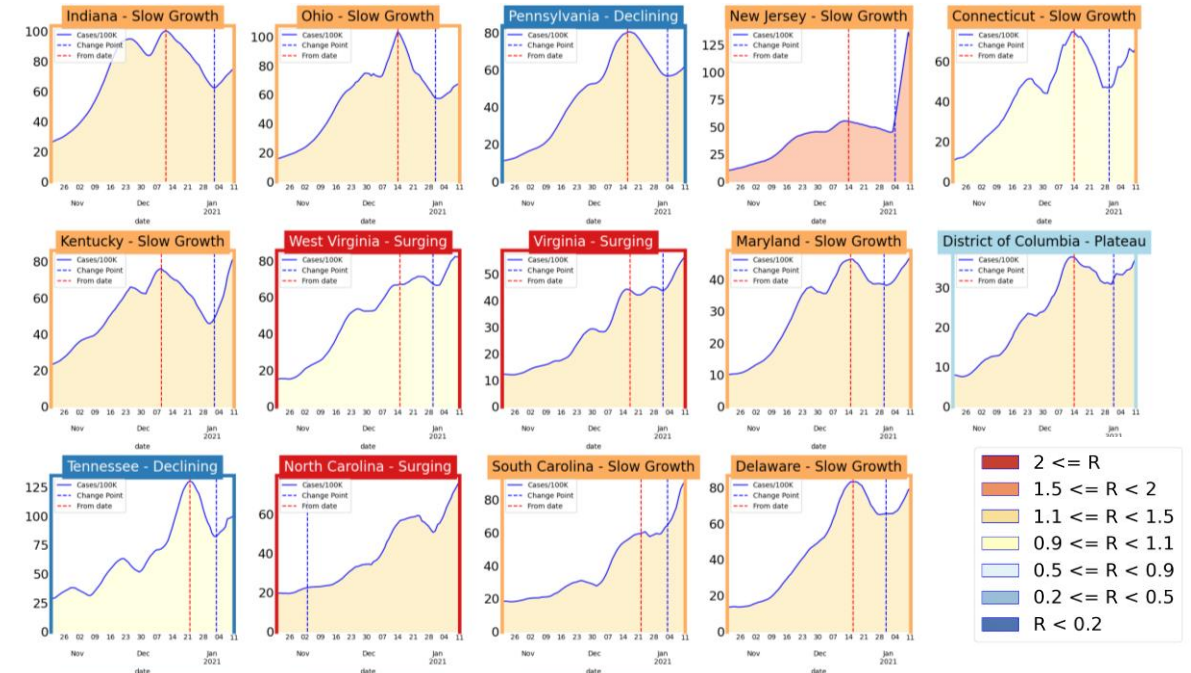
# Other State Comparisons

## Trajectories of States



- After several weeks of decline many states are rebounding
- Most Southern states are surging

## Virginia and her neighbors



- VA has rebounded and entered surging growth
- VA and her neighbors, mirror the patterns across the nation
- All neighbors are growing though some remain lower relative recent very high peaks

# Zip code level weekly Case Rate (per 100K)

## Case Rates in the last week by zip code

- Several of the top ten zip codes are home to prisons,
- Some counts are low and suppressed to protect anonymity, those are shown in white

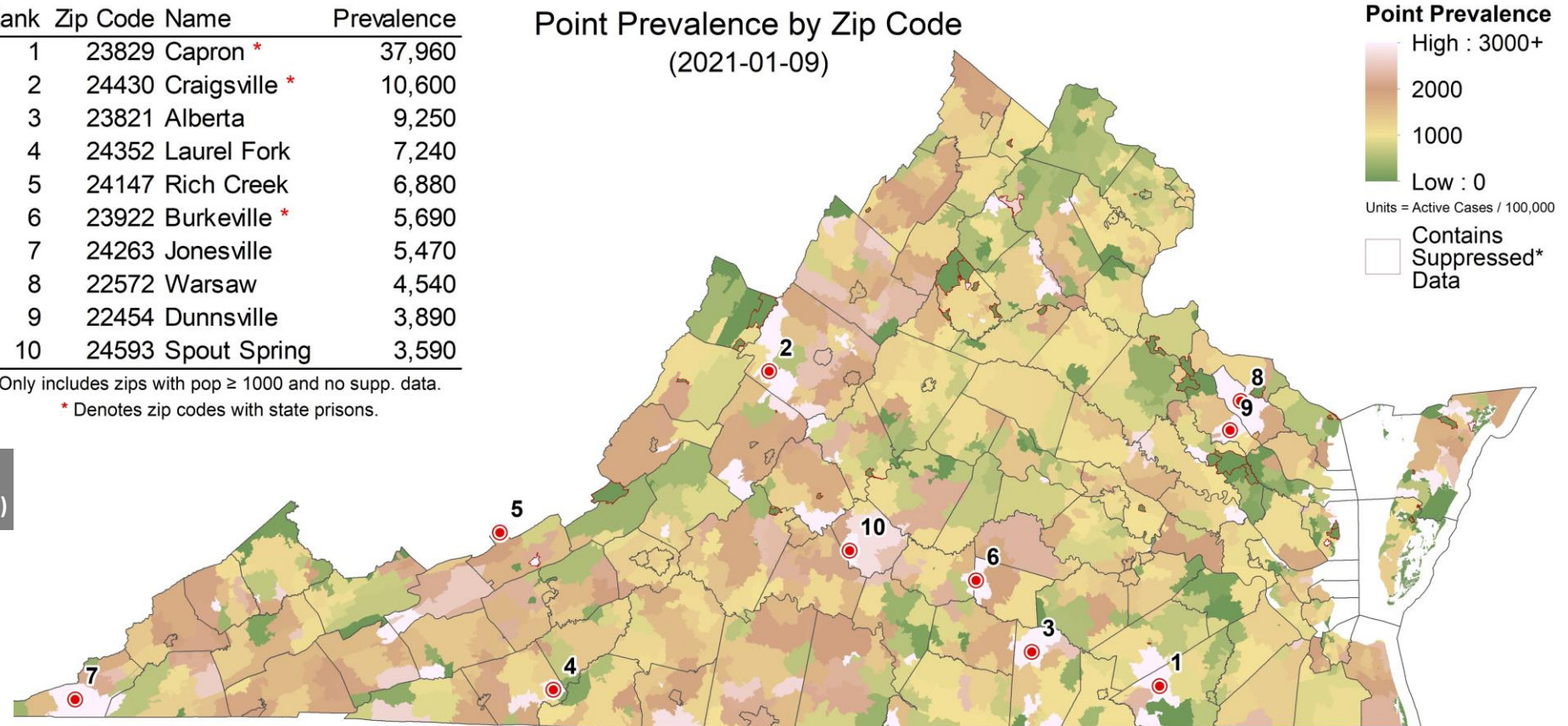
Rank	Zip Code	Name	Prevalence
1	23829	Capron *	37,960
2	24430	Craigsville *	10,600
3	23821	Alberta	9,250
4	24352	Laurel Fork	7,240
5	24147	Rich Creek	6,880
6	23922	Burkeville *	5,690
7	24263	Jonesville	5,470
8	22572	Warsaw	4,540
9	22454	Dunnsville	3,890
10	24593	Spout Spring	3,590

Only includes zips with pop ≥ 1000 and no supp. data.

\* Denotes zip codes with state prisons.

## Top 10 excluding prisons

Zip code	Name	Weekly Case Rate (per 100K)
23821	Alberta	9248
24352	Laurel Fork	7241
24147	Rich Creek	6883
24263	Jonesville	5475
22572	Warsaw	4536
22454	Dunnsville	3894
24593	Spout Spring	3590
23011	Barhamsville	3373
24083	Daleville	3329
22724	Jeffersonton	3251

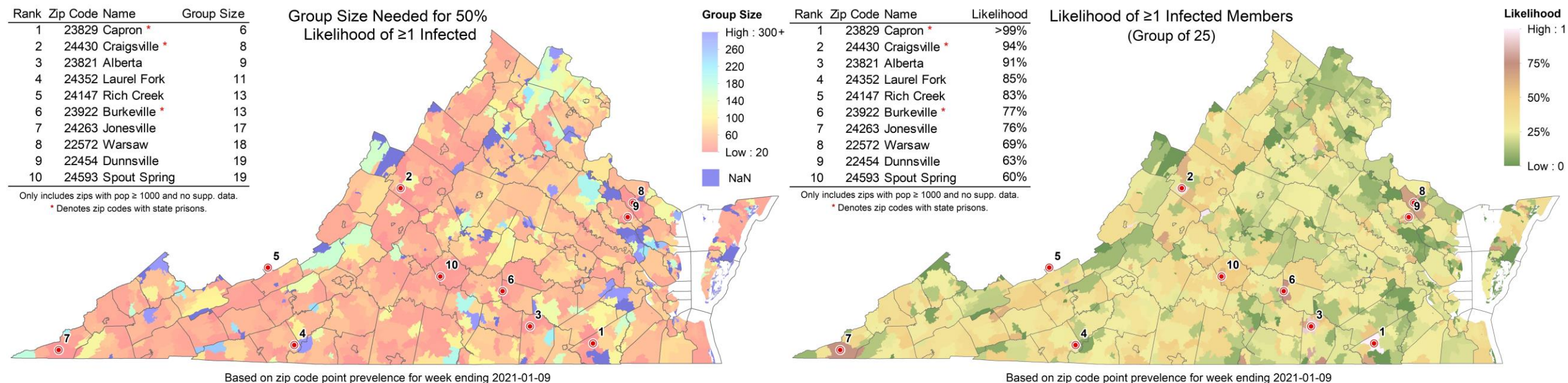




# Risk of Exposure by Group Size

## Case Prevalence in the last week by zip code used to calculate risk of encountering someone infected in a gathering of randomly selected people (group size 25)

- Assumes 3 undetected infections per confirmed case (ascertainment rate from recent seroprevalence survey)
- On left, minimum size of a group with a 50% chance an individual is infected by zip code (eg in a group of 20 in Staunton, there is a 50% chance someone will be infected)
- Some zip codes have high likelihood of exposure even in groups of 25



# New variants of SARS-CoV2

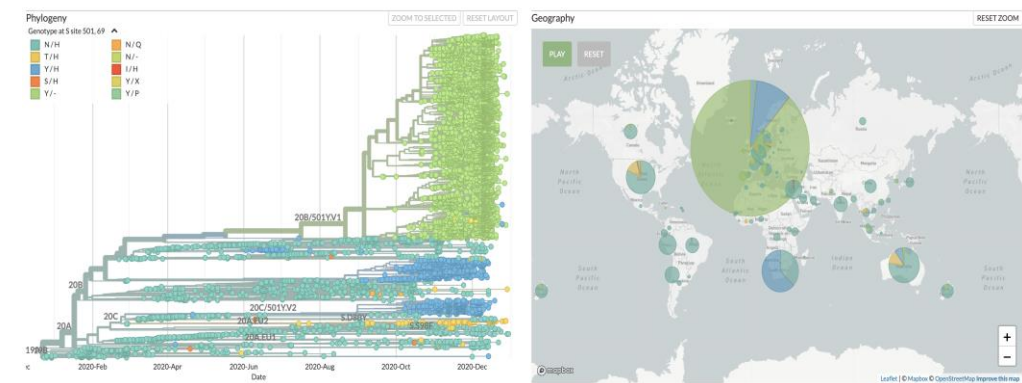
## Emerging new variant with increased transmissibility but no evidence of higher severity

- Aliases: [Variant VUI 202012/01](#) and [Lineage B.1.1.7](#)
- Variant has been detected in 10 states; most without travel history indicating local transmission
- This variant is still detected by PCR, and can be detected by proxy with an “S dropout”
- Evolution expected when virus under selective pressure
- Unlikely to alter efficacy of vaccines or other immune treatments
- NERVTAG suggests that “[VUI-202012/01 demonstrates a substantial increase in transmissibility compared to other variants](#)”
- Recent studies have been confirming that these variants aren’t likely to “escape” the current PCR tests nor approved vaccines

Phylogenetic analysis of SARS-CoV-2 clusters in their international context - cluster S.N501

Built with emmahodcroft/hcov\_cluster. Maintained by Emma Hodcroft and Richard Neher.

Showing 8013 of 8013 genomes sampled between Dec 2019 and Dec 2020.



[nextstrain.org](https://nextstrain.org)

# Model Update – Adaptive Fitting

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# Adaptive Fitting Approach

## Each county fit precisely, with recent trends used for future projection

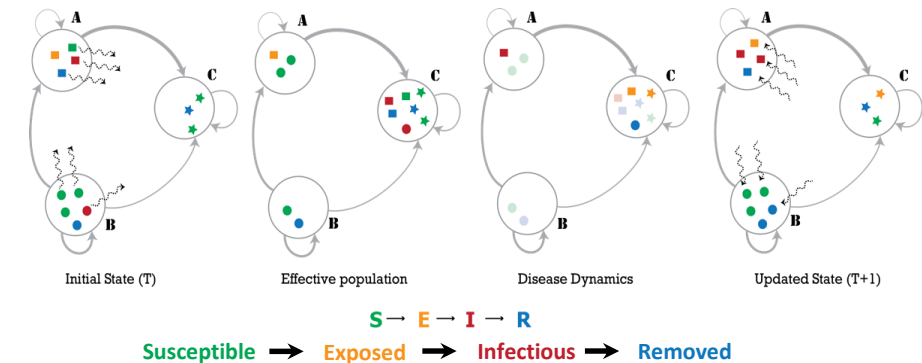
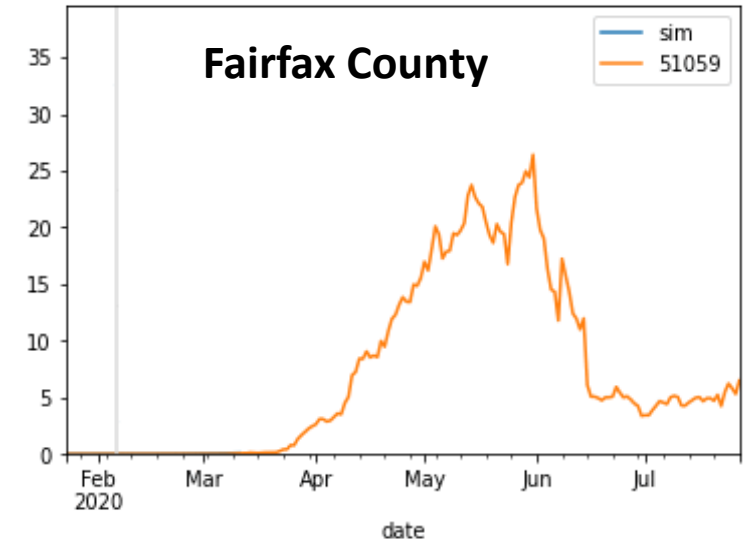
- Allows history to be precisely captured, and used to guide bounds on projections

## Model: An alternative use of the same meta-population model, PatchSim

- Allows for future “what-if” Scenarios to be layered on top of calibrated model
- Eliminates connectivity between patches, to allow calibration to capture the increasingly unsynchronized epidemic

## External Seeding: Steady low-level importation

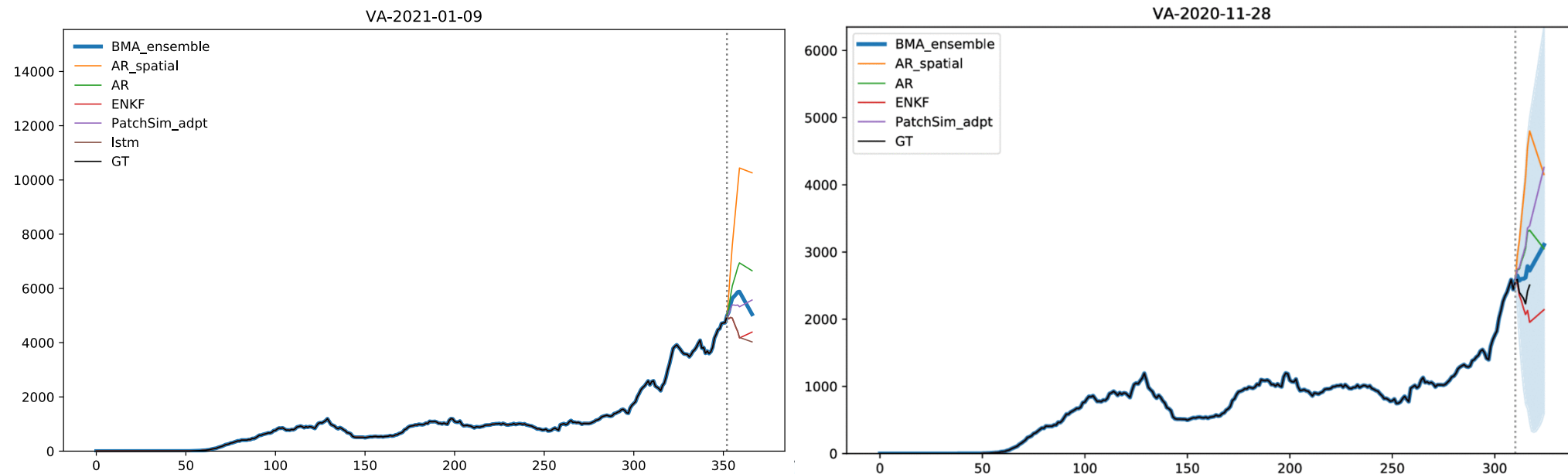
- Widespread pandemic eliminates sensitivity to initial conditions
- Uses steady 1 case per 10M population per day external seeding



# Using Ensemble Model to Guide Projections

An ensemble methodology that combines the Adaptive Fitting and machine learning and statistical models has been developed and refined

- **Models:** Adaptive Fitting, ARIMA, LSTM, AR, spatially driven AR, Kalman Filters (ENKF)
- This approach facilitates the use of other data streams (weather, mobility, etc.)
- Ensemble provides scaffolding for the Adaptive Fitting's short-term projections





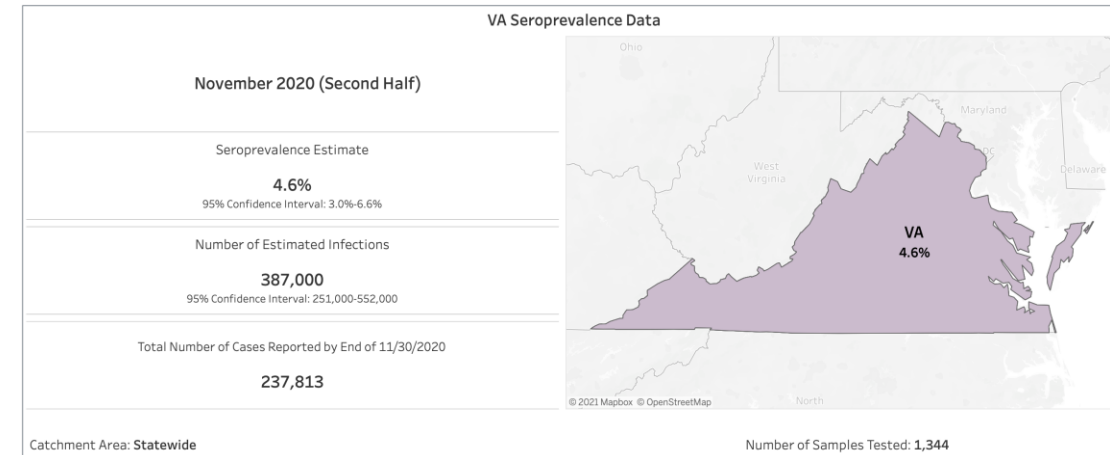
# Seroprevalence updates to model design

## Several seroprevalence studies provide better picture of how many actual infections have occurred

- Virginia Serology Study estimated 2.4% of Virginians estimated infected (as of Aug 15<sup>th</sup>)
- CDC Nationwide Commercial Laboratory Seroprevalence Survey estimated 4.6% [3.0% – 6.6%] seroprevalence as of Nov 12<sup>th</sup> – 26<sup>th</sup> up from 4.1% a month earlier

## These findings are equivalent to an ascertainment ratio of ~3x, with bounds of (1x to 7x)

- Thus for 3x there are 3 total infections in the population for every confirmed case
- Uncertainty design has been shifted to these bounds (previously higher ascertainment as was consistent earlier in the pandemic were being used)



<https://covid.cdc.gov/covid-data-tracker/#national-lab>

## Virginia Coronavirus Serology Project

Interim findings by region and statewide - July 22, 2020

Region	Number of participants	Number antibody positive	Crude prevalence per 100 participants	Weighted prevalence*	
				per 100 population	(95% CI)
Central	400	8	2.0	3.0	(0.5, 5.5)
East	707	9	1.3	1.5	(-0.2, 3.2)
Northern	819	36	4.4	4.2	(2.5, 5.9)
Northwest	756	11	1.5	0.9	(0.2, 1.6)
Southwest	431	3	0.7	1.0	(-0.2, 2.1)
Virginia	3,113	67	2.2	2.4	(1.6, 3.1)

\* Weighted prevalence is reweighted by region, age, sex, race, ethnicity, and insurance status to match census population.

<https://www.vdh.virginia.gov/content/uploads/sites/8/2020/08/VDH-Serology-Projects-Update-8-13-2020.pdf>

# Calibration Approach

- **Data:**
  - County level case counts by date of onset (from VDH)
  - Confirmed cases for model fitting
- **Calibration:** fit model to observed data and ensemble's forecast
  - Tune transmissibility across ranges of:
    - Duration of incubation (5-9 days), infectiousness (3-7 days)
    - Undocumented case rate (1x to 7x) guided by seroprevalence studies
    - Detection delay: exposure to confirmation (4-12 days)
  - Approach captures uncertainty, but allows model to precisely track the full trajectory of the outbreak
- **Project:** future cases and outcomes generated using the collection of fit models run into the future
  - **Mean trend from last 14 days of observed cases and first week of ensemble's forecast used**
  - Outliers removed based on variances in the previous 3 weeks
  - 2 week interpolation to smooth transitions in rapidly changing trajectories

## COVID-19 in Virginia:

Dashboard Updated: 1/13/2021  
Data entered by 5:00 PM the prior day.

Cases, Hospitalizations and Deaths					
Total Cases*		Total Hospitalizations**		Total Deaths	
<b>412,545</b>		<b>19,470</b>		<b>5,552</b>	
(New Cases: 4,598) <sup>^</sup>					
Confirmed†	Probable†	Confirmed†	Probable†	Confirmed†	Probable†
339,468	73,077	18,740	730	4,892	660

\* Includes both people with a positive test (Confirmed), and symptomatic with a known exposure to COVID-19 (Probable).  
 \*\* Hospitalization of a case is captured at the time VDH performs case investigation. This underrepresents the total number of hospitalizations in Virginia.  
<sup>^</sup>New cases represent the number of confirmed and probable cases reported to VDH in the past 24 hours.  
 † VDH adopted the updated CDC COVID-19 confirmed and probable surveillance case definitions on August 27, 2020. Found here: <https://www.cdc.gov/mmwr/conditions/coronavirus-disease-2019-covid-19/case-definition/20200805.html>

Outbreaks	
Total Outbreaks*	Outbreak Associated Cases
<b>2,088</b>	<b>49,621</b>

\* At least two (2) lab confirmed cases are required to classify an outbreak.

Testing (PCR Only)	
Testing Encounters PCR Only*	Current 7-Day Positivity Rate PCR Only**
<b>4,674,630</b>	<b>15.9%</b>

\* PCR\* refers to "Reverse transcriptase polymerase chain reaction laboratory testing."  
 \*\* Lab reports may not have been received yet. Percent positivity is not calculated for days with incomplete data.

Multisystem Inflammatory Syndrome in Children	
Total Cases*	Total Deaths
<b>13</b>	<b>0</b>

\*Cases defined by CDC HAN case definition: <https://emergency.cdc.gov/han/2020/han00437.asp>

Accessed 8:30am January 13, 2021

<https://www.vdh.virginia.gov/coronavirus/>

# Scenarios – Seasonal Effects

- Variety of factors continue to drive transmission rates
  - Seasonal impact of weather patterns, new variants of the virus with enhanced transmissibility, travel and gatherings related to holidays, fatigue with infection control practices
- Plausible levels of transmission can be bounded by past experience
  - Assess transmission levels at the county level since May 2020
  - Use the highest and lowest levels experienced (excluding outliers) as plausible bounds for levels of control achievable
  - Transition from current levels of projection to the new levels over 2 months
- New planning Scenarios:
  - **Best of the Past:** Lowest level of transmission (5th percentile)
  - **Fatigued Control:** Highest level of transmission (95<sup>th</sup> percentile) increased by additional 5%

# Scenarios – Vaccines

- Vaccination has started, and efforts are underway to increase its pace
  - Exact achievable rollouts and level of coverage are unknown
- Vaccine efficacy varies over course of vaccine
  - FDA EUAs show 50% efficacy achieved 2 weeks after first dose, and 95% two weeks after second dose
  - Assuming 3.5 week (average of Pfizer and Moderna) gap between doses
- Schedules
  - **Optimistic:** 25M courses in US (~ 660K in VA) starting in January, and continued 25M (~660K) per month,
  - **Pessimistic:** 12.5M people in US (~330K in VA) in January, then 25M (~660K) per month, with vaccine hesitancy leaving 50% of delivered vaccines unused.
  - Assume that vaccinations before January protect individuals but are not influencing transmission dynamics

Current rollouts and scenarios inspired by MIDAS Network COVID-19 Scenario Hub: <https://github.com/midas-network/covid19-scenario-modeling-hub>

# Scenarios – Seasonal Effects and Vaccines

Three scenarios combine these seasonal effects and vaccine scenarios

- **Adaptive:** No seasonal effects from base projection paired with optimistic vaccine schedule
- **Adaptive-FatigueControl:** Fatigued control seasonal effects paired with pessimistic vaccine schedule
- **Adaptive-BestPastControl:** Best of the past control seasonal effects paired with optimistic vaccine schedule
- Counterfactuals with no vaccine (“NoVax”) are provided for comparison purposes

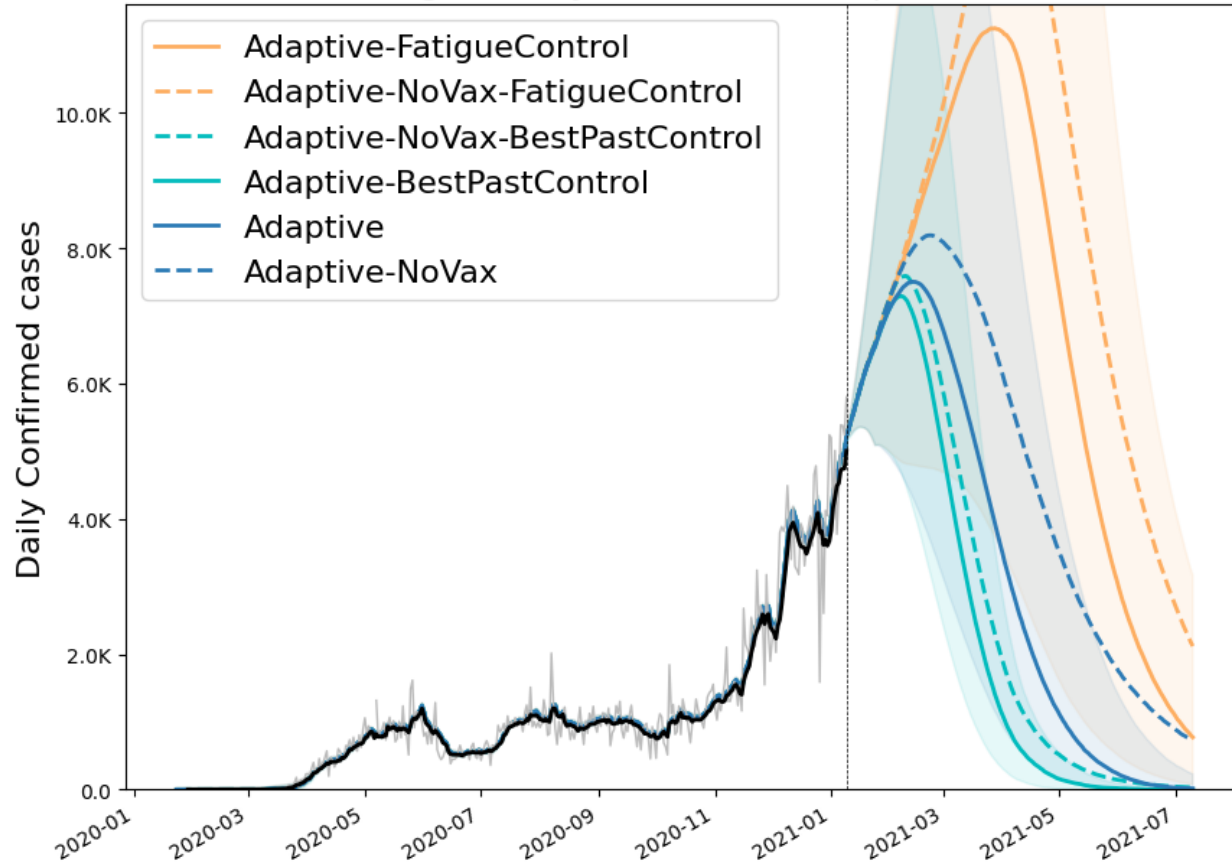
# Model Results

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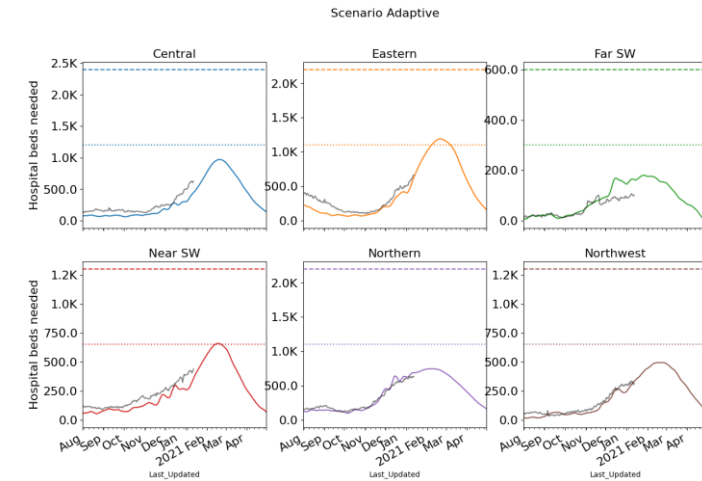
# Outcome Projections

## Confirmed cases

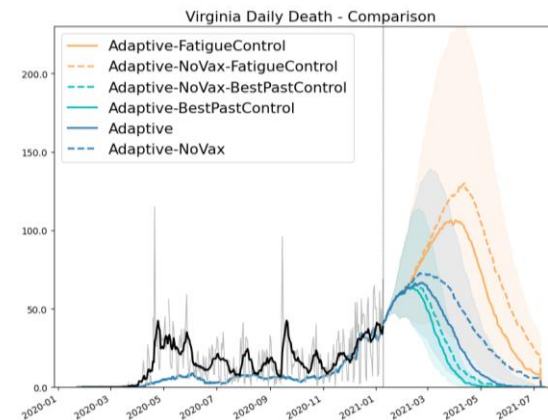
Virginia Daily Confirmed - Comparison



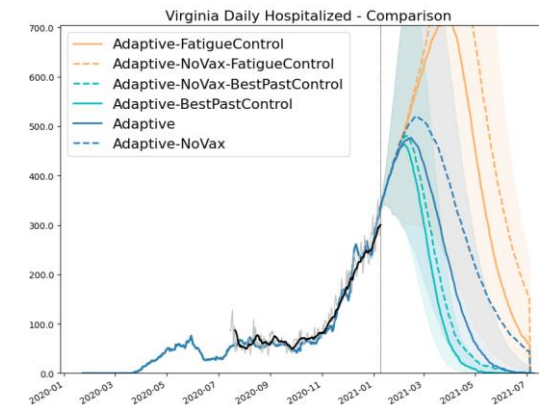
## Estimated Hospital Occupancy



## Daily Deaths



## Daily Hospitalized

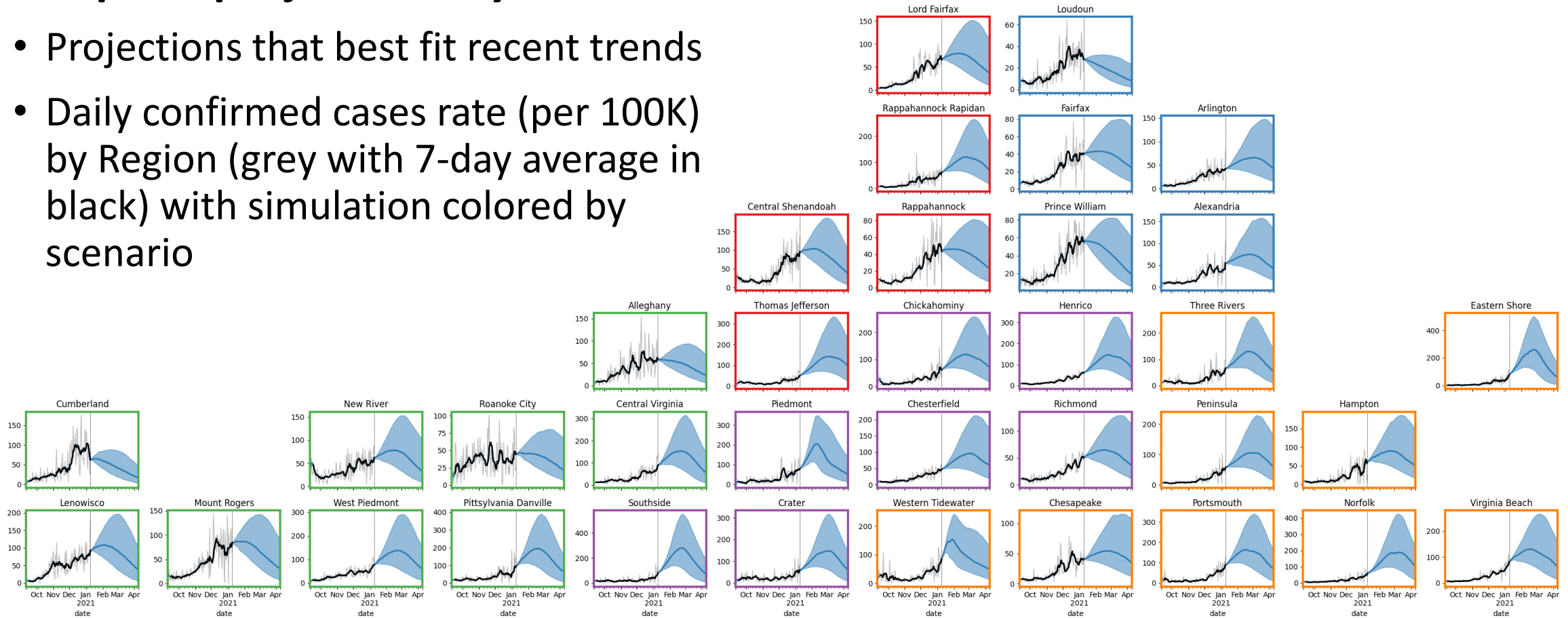




# District Level Projections: Adaptive

## Adaptive projections by District

- Projections that best fit recent trends
- Daily confirmed cases rate (per 100K) by Region (grey with 7-day average in black) with simulation colored by scenario

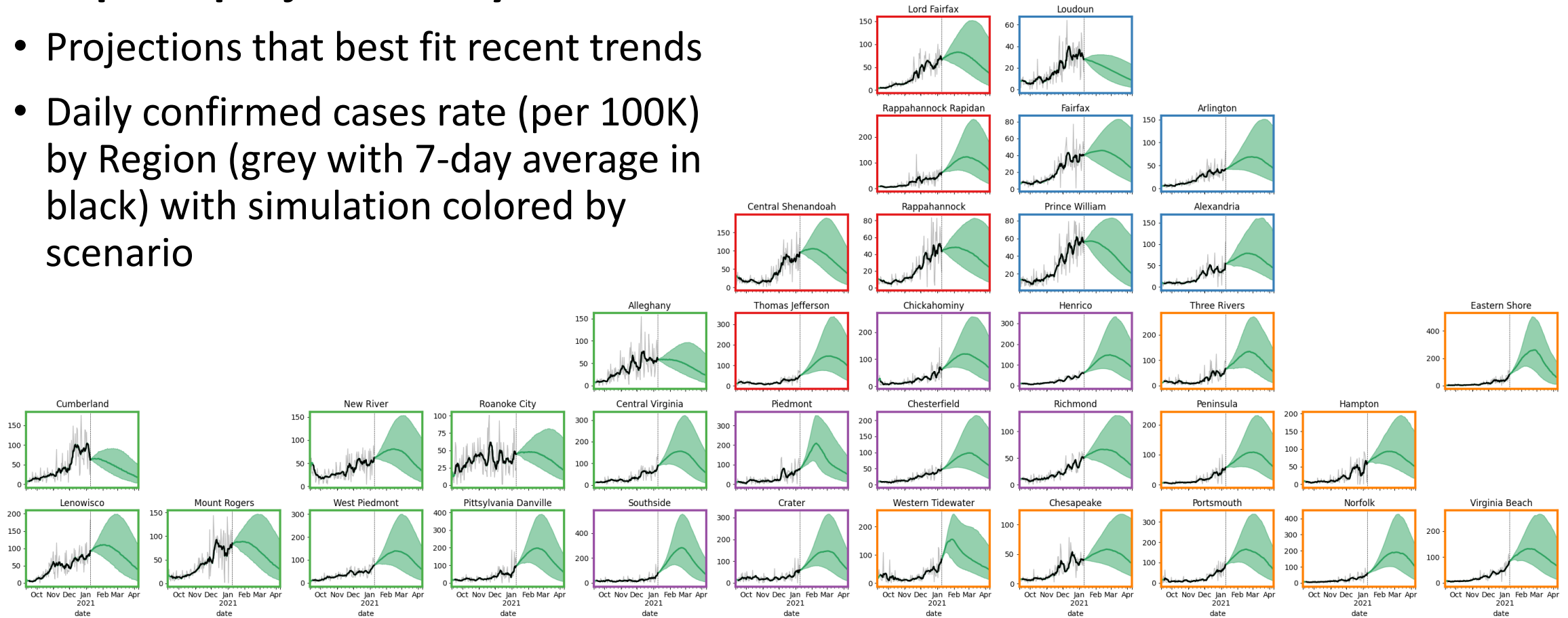




# District Level Projections: Adaptive-BestPastControl

## Adaptive projections by District

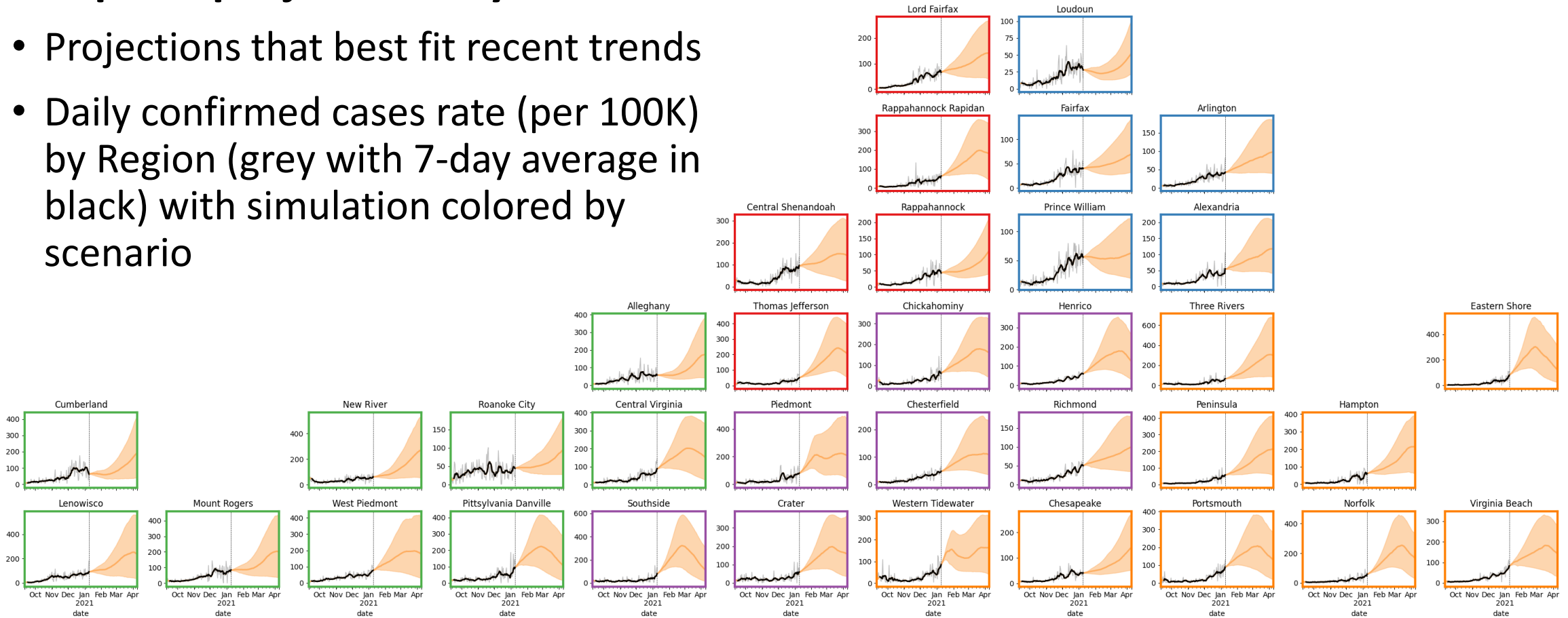
- Projections that best fit recent trends
- Daily confirmed cases rate (per 100K) by Region (grey with 7-day average in black) with simulation colored by scenario



# District Level Projections: Adaptive-FatigueControl

## Adaptive projections by District

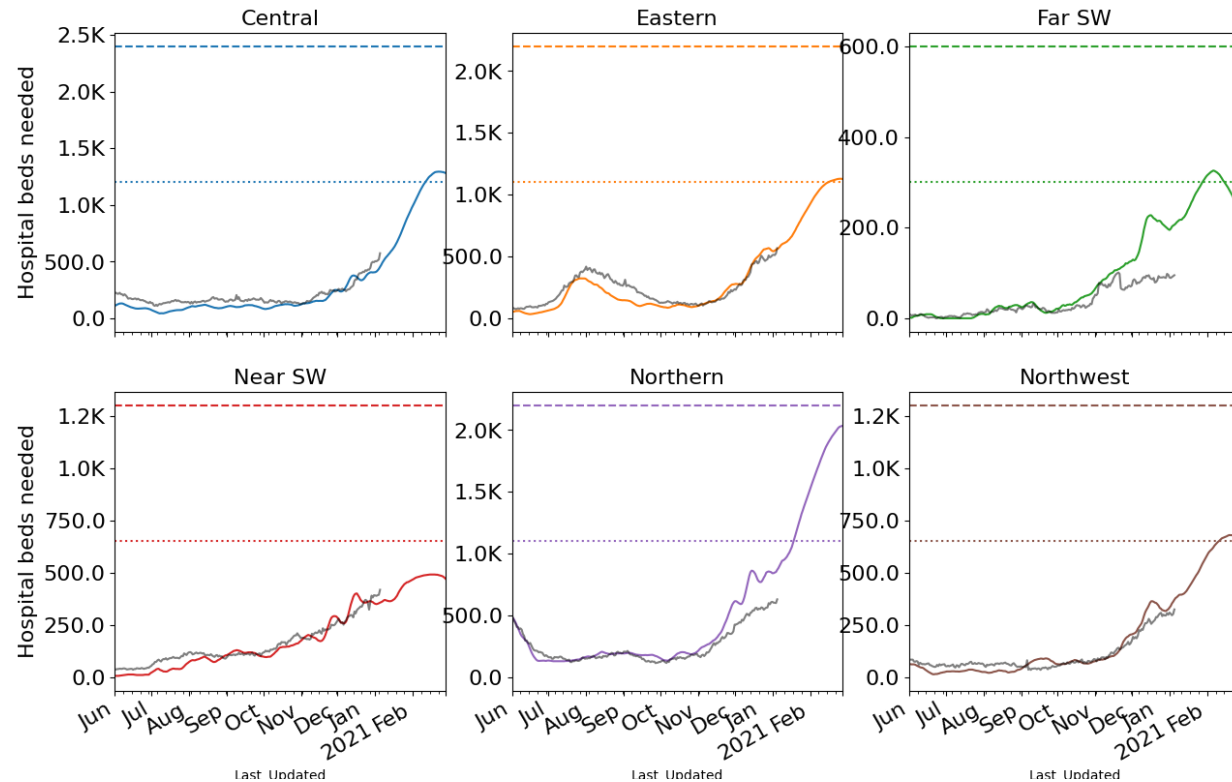
- Projections that best fit recent trends
- Daily confirmed cases rate (per 100K) by Region (grey with 7-day average in black) with simulation colored by scenario



# Hospital Demand and Bed Capacity by Region

## Capacities\* by Region – Adaptive-FatigueControl

COVID-19 capacity ranges from 80% (dots) to 120% (dash) of total beds



Week Ending	Adaptive	Adaptive-FatigueControl
1/10/21	33,297	33,298
1/17/21	38,230	38,234
1/24/21	42,524	42,580
1/31/21	46,142	46,494
2/7/21	48,872	50,207
2/14/21	50,232	54,228
2/21/21	49,776	58,589
2/28/21	47,436	62,628
3/7/21	43,482	66,818
3/14/21	38,870	70,870
3/21/21	33,802	73,814
3/28/21	28,472	75,178

Weekly confirmed cases

## If Adaptive-FatigueControl scenario persists:

- All regions approach initial bed capacity this winter
- Surge capacity approached in Northern region in Feb to early March

\* Assumes average length of stay of 8 days

13-Jan-21

# Key Takeaways

Projecting future cases precisely is impossible and unnecessary.  
Even without perfect projections, we can confidently draw conclusions:

- **Case rate growth in Virginia continues to surge along with the nation, but remains below national average**
- VA mean weekly incidence (60/100K) up (from 52) as national levels also rebound (to 67/100K from 60/100K); Virginia records highest daily case rate in past week
- Projections are mostly up across commonwealth
- Recent updates:
  - Modified scenarios to be based on past control levels (best and fatigued)
  - Refined vaccination schedule to account for partial protection from first dose
  - Planning scenarios start changing transmission rates at end of the month (Jan 30)
  - Adjusted “rescaling” method and data sources to better accommodate recent trends
- The situation is changing rapidly. Models will be updated regularly.

# References

Venkatramanan, S., et al. "Optimizing spatial allocation of seasonal influenza vaccine under temporal constraints." *PLoS computational biology* 15.9 (2019): e1007111.

Arindam Fadikar, Dave Higdon, Jiangzhuo Chen, Bryan Lewis, Srinivasan Venkatramanan, and Madhav Marathe. Calibrating a stochastic, agent-based model using quantile-based emulation. *SIAM/ASA Journal on Uncertainty Quantification*, 6(4):1685–1706, 2018.

Adiga, Aniruddha, Srinivasan Venkatramanan, Akhil Peddireddy, et al. "Evaluating the impact of international airline suspensions on COVID-19 direct importation risk." *medRxiv* (2020)

NSSAC. PatchSim: Code for simulating the metapopulation SEIR model. <https://github.com/NSSAC/PatchSim> (Accessed on 04/10/2020).

Virginia Department of Health. COVID-19 in Virginia. <http://www.vdh.virginia.gov/coronavirus/> (Accessed on 04/10/2020)

Biocomplexity Institute. COVID-19 Surveillance Dashboard. <https://nssac.bii.virginia.edu/covid-19/dashboard/>

Google. COVID-19 community mobility reports. <https://www.google.com/covid19/mobility/>

Biocomplexity page for data and other resources related to COVID-19: <https://covid19.biocomplexity.virginia.edu/>

# Questions?

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# Supplemental Slides

# Estimating Daily Reproductive Number

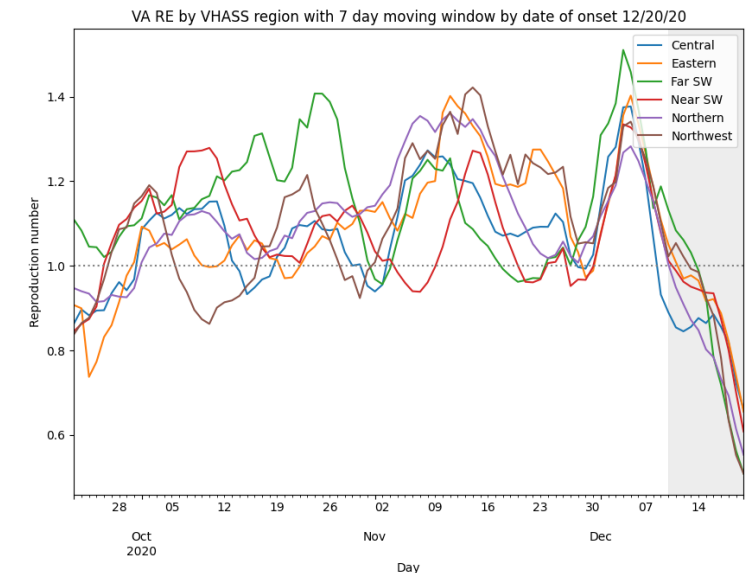
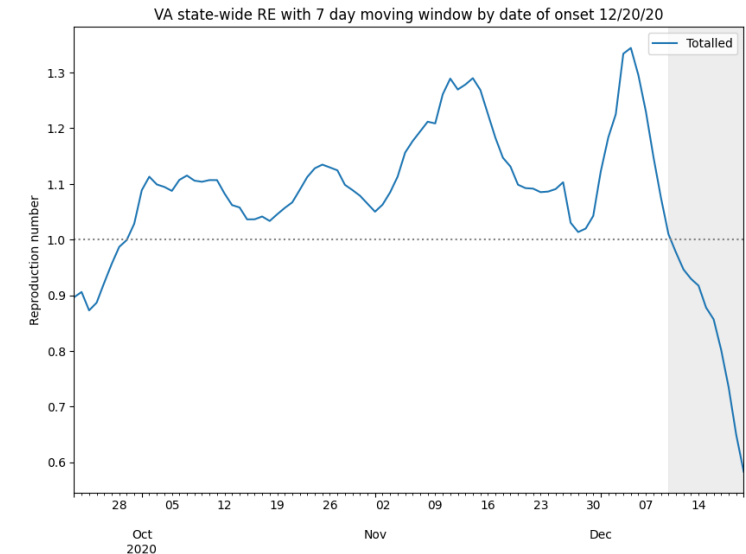
## Dec 11<sup>th</sup> Estimates

Region	Date of Onset $R_e$	Date Onset Diff Last Week
State-wide	0.977	-0.275
Central	0.854	-0.491
Eastern	1.009	-0.176
Far SW	1.083	-0.323
Near SW	0.988	-0.301
Northern	0.949	-0.253
Northwest	1.054	-0.193

### Methodology

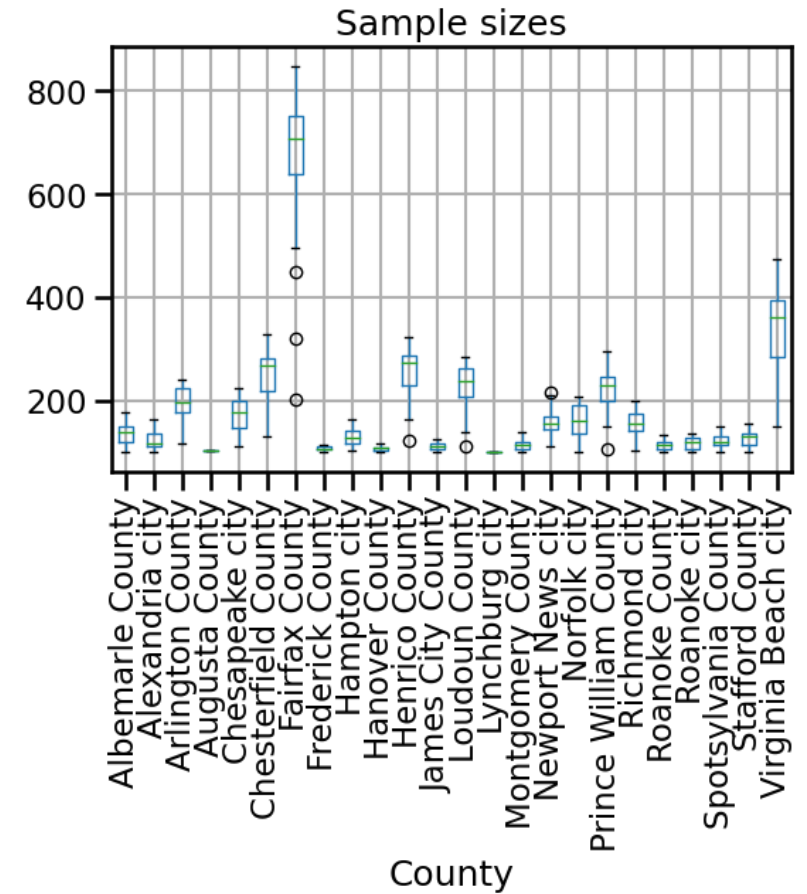
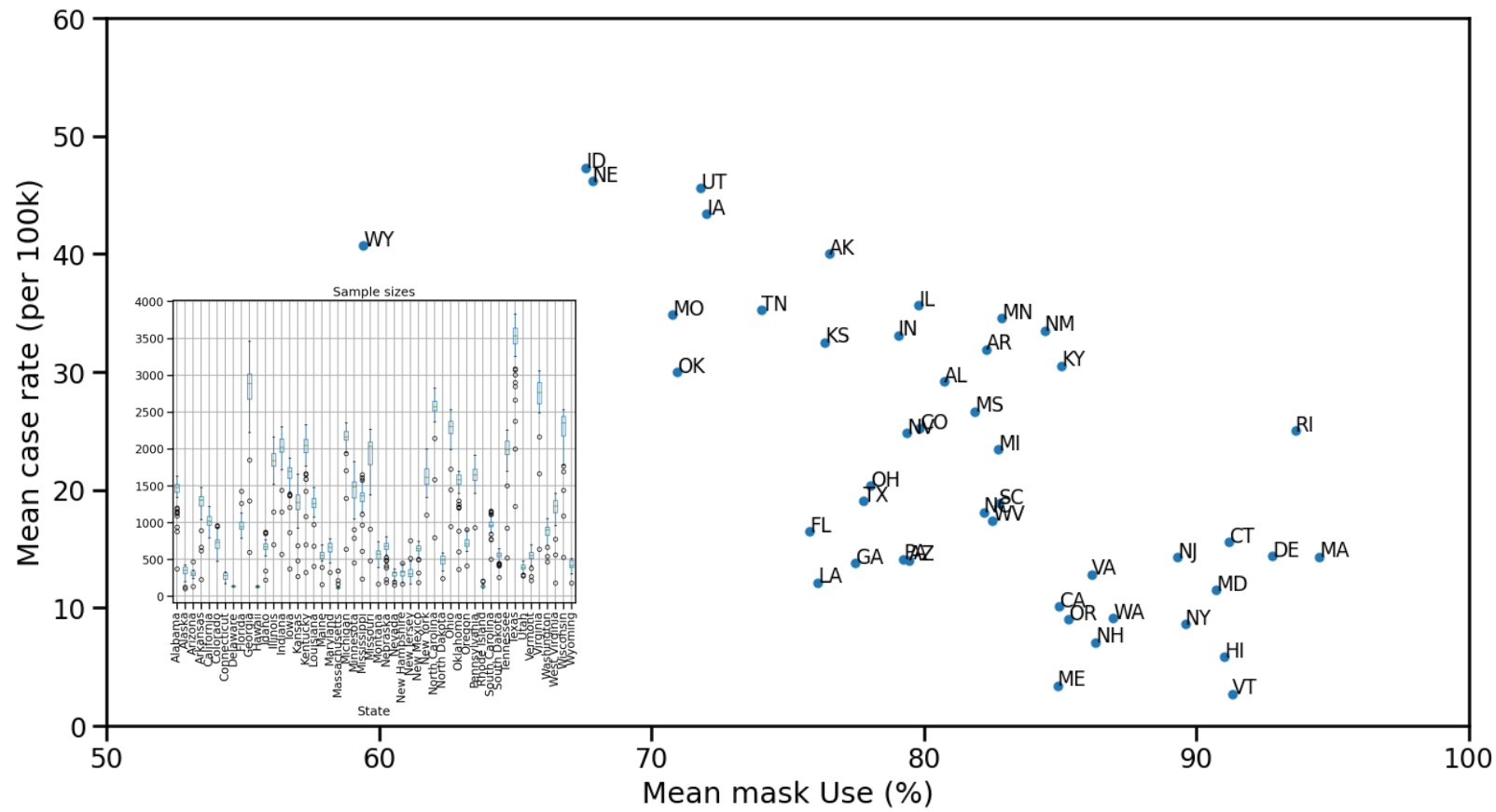
- Wallinga-Teunis method (EpiEstim<sup>1</sup>) for cases by date of onset
- Serial interval: 6 days (2 day std dev)
- Recent estimates may be unstable due to backfill

1. Anne Cori, Neil M. Ferguson, Christophe Fraser, Simon Cauchemez. A New Framework and Software to Estimate Time-Varying Reproduction Numbers During Epidemics. American Journal of Epidemiology, Volume 178, Issue 9, 1 November 2013, Pages 1505–1512, <https://doi.org/10.1093/aje/kwt133>





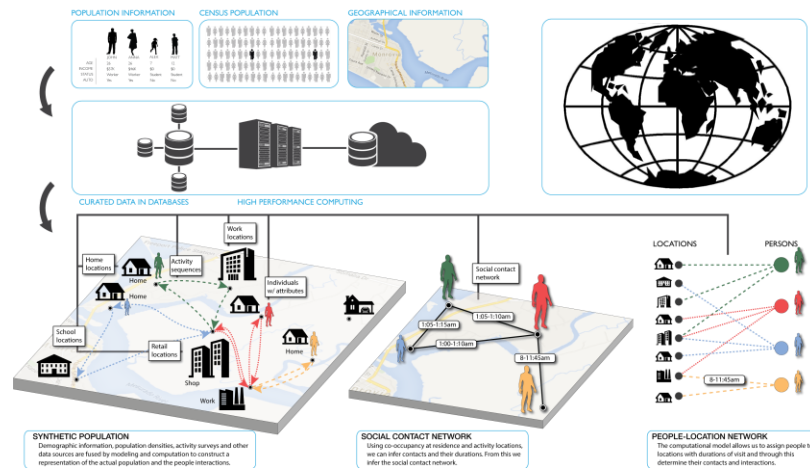
# Mask usage sample sizes



# Agent-based Model (ABM )

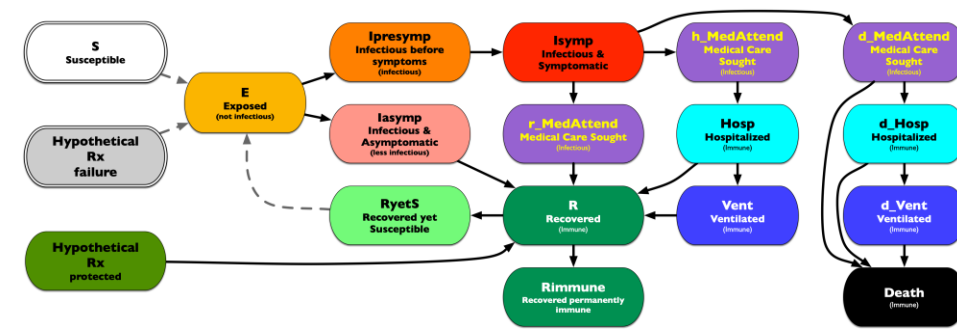
## EpiHiper: Distributed network-based stochastic disease transmission simulations

- Assess the impact on transmission under different conditions
- Assess the impacts of contact tracing



### Synthetic Population

- Census derived age and household structure
- Time-Use survey driven activities at appropriate locations



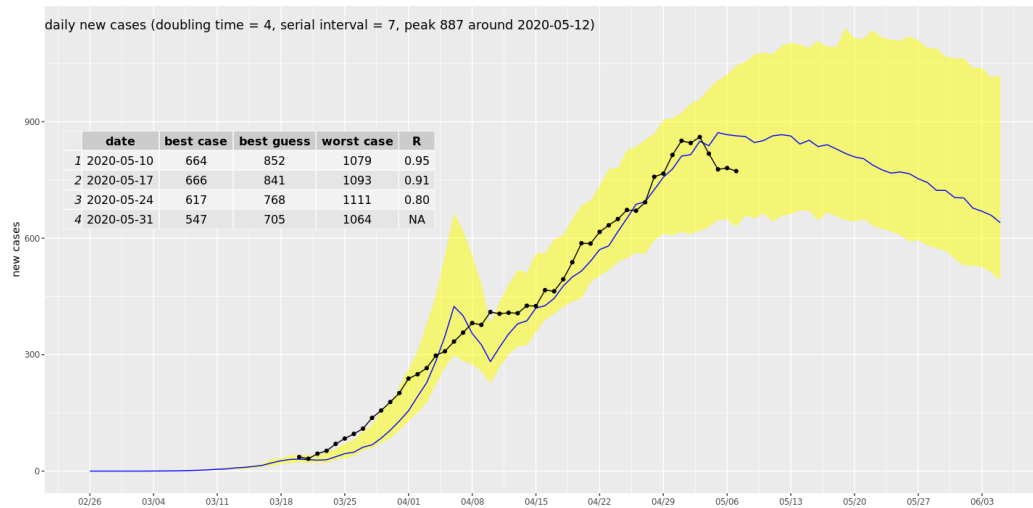
### Detailed Disease Course of COVID-19

- Literature based probabilities of outcomes with appropriate delays
- Varying levels of infectiousness
- Hypothetical treatments for future developments

# ABM Social Distancing Rebound Study Design

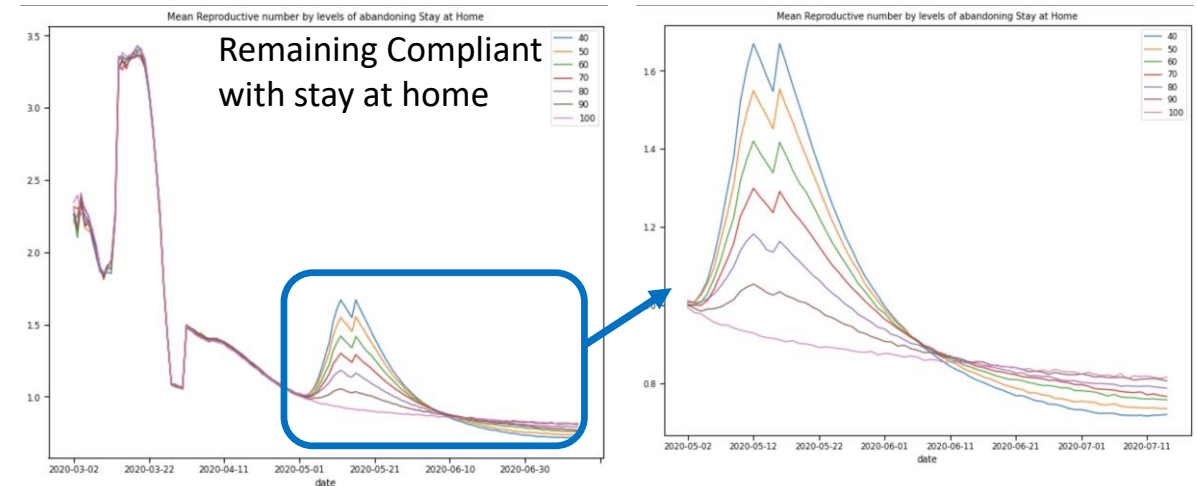
## Study of "Stay Home" policy adherence

- Calibration to current state in epidemic
- Implement "release" of different proportions of people from "staying at home"



### Calibration to Current State

- Adjust transmission and adherence to current policies to current observations
- For Virginia, with same seeding approach as PatchSim



### Impacts on Reproductive number with release

- After release, spike in transmission driven by additional interactions at work, retail, and other
- At 25% release (70-80% remain compliant)
- Translates to 15% increase in transmission, which represents a  $1/6^{\text{th}}$  return to pre-pandemic levels

# Medical Resource Demand Dashboard

<https://nssac.bii.virginia.edu/covid-19/vmrddash/>

